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           CHALUP, Michael S.
          CHANG, Simon C.
CHEN, Alice
D'SA, Steven A.
           AMSHEY, Stefan
           DAHL, Christopher R.
           DAM, Tam C.
           DANIELS, Susan E.
          DUFOUR, Gerard E.
FLORES, Vincent
FONG, Willy T.
GREENAWALT, Lila B.
           HILLMAN, Jennifer L.
           JONES, Anissa L.
           LIU, Tommy F.
           ROSEBERRY, Ann M.
           ROSEN, Bruce H. RUSSO, Frank D.
           STOCKDREHER, Theresa K.
           DAFFO, Abel
WRIGHT, Rachel J.
           YAP, Pierre E.
YU, Jimmy Y.
           BRADLEY, Diana L. BRATCHER, Shawn R.
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agecegagaa atgacegace tgatgegtgt tataacecat etgageeege tgeaaceace 360
gtttttgaaa taagaactgg ccttttgcta ggtgatacaa gtaaaataat aactagaaca 420 gaagaaaaag aaatccccaa gcaaataact ttaagatttg atgcttgtgc agccattaat 480
aqtaaaaaqc tagaaatagg atqtqqttct cttaactgag aaaggagcta aagagtagaa 540
aataaatatg tttgtcatga gtcag
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<210> 209
<211> 567
<212> DNA
<213> Homo sapiens
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acagttcaca cttactactg ttgtagttat aaattattca cttgttttaa ataagcacat 60
atatgatggc tgagtgagaa tgatctagaa ctatgccagg gattagataa aaacacagag 120 taaaagttgg ccatatagta acattttga tgtccaaatt ggagaggaat ccctcaaatg 180
ttcaaaaata taggaatggg tcgggtgtgg tggctcatgc ctgtaatccc agcactttgg 240
gaggecaagg caggaagate acttgageee aggagtteaa gaccageetg ggcaacacag 300
tgagaccctg actcaaaaaa taaaaagaaa agacaaaaat acaagaatgg ctaagcaaac 360
tagigtacai caacciggag gaatattata tagictitta aaataacaaa taigtagatt 420
gttaatacat gggaaaaggt atatgaaata ggtggaacca agcagaactg agtcccaagg 480 tgggtgggaa agaggctgcc atgctggctg aggtacctga gaacaatgaa aacaagtcat 540
attaagacca atatcaatat cccactg
<210> 210
<211> 971
<212> DNA
<213> Homo sapiens
<221> misc_feature
<223> Incyte ID No: LG:380497.2:2000MAY19
tegeceegte egetgeagte egeeggegag ggagttaege aegteetgat teteetggag 60 tetecageee geecagtgge egeagteace eaggteeaga ggeggeggta teacaggete 120
tecgacatgt ctatgetgge tgaaegtegg eggaageaga agtgggetgt ggateeteag 180
aacactgcct ggagtaatga cgattccaag tttggccagc ggatgctaga gaagatgggg 240 tggtctaaag gaaagggttt aggggctcag gagcaaggag ccacagatca tattaaagtt 300
caagtgaaaa ataaccacct gggactcgga gctaccatca ataatgaaga caactggatt 360
gcccatcagg atgattttaa ccagettetg gccgaactga acaettgcca tgggcaggaa 420 accacagatt cctcggacaa gaaggaaaag aaatetttta gccttgagga aaagtccaaa 480
atctccaaaa accgtgttca ctatatgaaa ttcacaaaag ggcgatgcca gtccctccac 540
tccagaggag aacgaaacca cgacaaccag cgccttcacc atccaggagt actttgccaa 600
geggatggea geactgaaga acaageeeea ggtteeagtt ceagggtetg acatttetga 660 gaegeaggtg gaacgtaaaa gggggaagaa aagaaataaa gaggeeacag gtaaagatgt 720
ggaaagttac ctccagccta aggccaagag gcacacggag ggaaagcccg agagggccga 780
ggcccaggag cgagtggcca agaagaagag cgcgccagca gaagagcagc tcagaggccc 840 ctgctgggac cagagttcca aggcctctgc tcaggatgca ggggaccatg tgcagccgcc 900
tgagggccgg gacttcaccc tgaaagccaa aaagaggaga ggcgagaaaa agctgcaaaa 960
accagtagag a
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<211> 486
<212> DNA
<213> Homo sapiens
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<221> misc_feature
<223> Incyte ID No: LI:272913.22:2000MAY01
<400> 211
agegegacag eggeageace aagggaaegg aaaatggege eteaeggeee gggtagtett 60
acgaecetgg tgeeetggge tgeegeeetg eteetegete tgggegtgga aagggetetg 120
gegetaceeg agatatgeac ceaatgteea gggagegtge aaaatttgte aaaagtggee 180
ttttattgta aaacgacacg agagetaatg etgeatgeee gttgetgeet gaateagaag 240 gaeeetggte caaactttea teaggeacat accaetgtea teatagacet geaageaaac 300
cccctcaaag gtgacttggc caacacttc cgtggcttta ctcagctcca gactctgata 360
ctgccacaac atgtcaactg ttctggagga attaatgcct ggaatactat cacctcttat 420 atagacaacc aaatctgtca agggcaaaag aacctttgca ataacactgg ggacccagaa 480
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atgtgt <210> 212 <211> 115 <212> PRT <213> Homo sapiens <220> <221> misc\_feature <223> Incyte ID No: LG:1040582.1.orf3:2000FEB18 <400> 212 Ile Leu Ala Thr Leu Ser Ser Pro Gln Pro Val Leu Ile Ala Lys 10 Gly Thr Met Thr Ala Ser Ser Val Leu Leu His Thr Gly Gln Lys 20 25 Met Pro Leu Ile Gly Leu Gly Thr Trp Lys Ser Glu Pro Gly Gln Val Lys Ala Ala Ile Lys Tyr Ala Leu Ser Val Gly Tyr Arg His 50 55 60 Ile Asp Cys Ala Ser Val Tyr Gly Asn Glu Thr Glu Ile Gly Glu 75 65 70 Ala Leu Lys Glu Ser Val Gly Ser Gly Lys Ala Val Pro Arg Glu 90 80 85 Glu Leu Phe Val Thr. Ser Lys Leu Trp Asn Thr Lys His His Pro 95 100 105 Glu Asp Val Glu Pro Ala Leu Arg Lys Thr 110 <210> 213 <211> 161 <212> PRT <213> Homo sapiens <220> <221> misc\_feature <223> Incyte ID No: LG:453570.1.orf3:2000FEB18 <400> 213 Cys Thr Leu Leu Arg Leu Ile Pro Ser Ala Ser His Phe Lys Arg 10 15 1 Phe Asp Arg Val Arg Arg Phe Ala Pro Ala Ala Met Ala Thr Ser 25 30 20 Ser Gly Pro Lys Glu Ala Pro Ala Asn Asn Pro Gly Leu Gln Thr 45 40 Glu Val Asp Pro Ala Thr Lys Gly Tyr Phe Leu Gln Gln Thr Met 50 55 60 Leu Arg Val Lys Asp Pro Lys Val Ser Leu Asp Phe Tyr Ser Arg 65 70 Val Met Gly Met Ser Leu Leu Lys Arg Leu Asp Phe Glu Glu Met 90 80 85 Lys Phe Ser Leu Tyr Phe Leu Gly Tyr Glu Asp Val Thr Leu Ala 95 100 105 Pro Asp Asp His Ile Lys Arg Thr Glu Trp Thr Phe Arg Gln Lys 110 115 Ala Thr Leu Glu Leu Thr His Asn Trp Gly Thr Glu Asn Asp Pro 130 135 125 Glu Phe Lys Gly Tyr His Asn Gly Asn Ser Asp Pro Arg Gly Phe 140 145 150 Gly His Ile Gly Val Thr Val Asp Asp Val His 155 160 <210> 214 <211> 332 <212> PRT

<213> Homo sapiens

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<223> Incyte ID No: LG:408751.3.orf2:2000FEB18
Arg Asp Pro Gly Trp Gln Ile Arg Asp Arg Ala Gly Leu Ala Trp
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                                                           15
Asn Met Leu Ala Asn Ser Ala Ser Val Arg
                                        Ile Leu Ile Lys Gly
                                                           30
                 20
Gly Lys Val Val Asn Asp Asp Cys Thr His Glu Ala Asp Val
                                                           45
                 35
                                      40
Ile Glu Asn Gly Ile Ile Gln Gln Val Gly Arg Glu Leu Met Ile
                 50
Pro Gly Gly Ala Lys Val Ile Asp Ala Thr Gly Lys Leu Val Ile
                                      70
                 65
Pro Gly Gly Ile Asp Thr Ser Thr His Phe His Gln Thr Phe Met
                 80
                                      85
                                                           90
Asn Ala Thr Cys Val Asp Asp Phe Tyr His Gly Thr Lys Ala Ala
                 95
                                     100
                                                          105
Leu Val Gly Gly Thr Thr Met Ile Ile Gly His Val Leu Pro Asp
                110
                                     115
                                                          120
Lys Glu Thr Ser Leu Val Asp Ala Tyr Glu Lys Cys Arg Gly Leu
                125
                                     130
                                                          135
Ala Asp Pro Lys Val Cys Cys Asp Tyr Ala Leu His Val Gly Ile
                                                          150
                                     145
                140
Thr Trp Trp Ala Pro Lys Val Lys Ala Glu Met Glu Thr Leu Val
                                                          165
                155
                                     160
Arg Glu Lys Gly Val Asn Ser Phe Gln Met Phe Met Thr Tyr Lys
                170
                                     175
                                                          180
Asp Leu Tyr Met Leu Arg Asp Ser Glu Leu Tyr Gln Val Leu His
                185
                                     190
                                                          195
Ala Cys Lys Asp Ile Gly Ala Ile Ala Arg Val His Ala Glu Asn
                                     205
                200
                                                          210
Gly Glu Leu Val Ala Glu Gly Ala Lys Glu Ala Leu Asp Leu Gly
                                     220
                                                          225
                215
Ile Thr Gly Pro Glu Gly Ile Glu Ile Ser Arg Pro Glu Glu Leu
                230
                                     235
Glu Ala Glu Ala Thr His Arg Val Ile Thr Ile Ala Asn Arg Thr
                 245
                                     250
                                                          255
His Cys Pro Ile Tyr Leu Val Asn Val Ser Ser Ile Ser Ala Gly
                                                          270
                260
                                     265
Asp Val Ile Ala Ala Ala Lys Met Gln Gly Lys Val Val Leu Ala
                                                          285
                275
                                     280
Glu Thr Thr Thr Ala His Ala Thr Leu Thr Gly Leu His Tyr
                                                         Tyr
                                     295
                                                          300
                290
His Gln Asp Trp Ser His Ala Ala Ala Tyr Val Thr Val Pro
                                                         Pro
                305
                                     310
Leu Arg Leu Asp Thr Asn Thr Ser Thr Tyr Leu Met Ser Leu Leu
                320
                                     325
                                                          330
Ala Lys
<210> 215
<211> 274
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte ID No: LI:090574.1.orf3:2000FEB01
<400> 215
Ala Arg Asn Ser Ala Arg Val Pro Pro Arg Gly Thr Met Ser Arg
                                      10
                                                           15
                                                         Lys
                Tyr Arg Glu His Asn Gly Pro Ile His Trp
                  20
Glu Phe Phe Pro Ile Ala Asp Gly Asp Gln Gln Ser Pro Ile Glu
```

```
35
                                      40
Ile Lys Thr Lys Glu Val Lys Tyr Asp Ser Ser Leu Arg Pro Leu
                 50
                                      55
                                                          60
Ser Ile Lys Tyr Asp Pro Ser Ser Ala Lys
                                        Ile Ile Ser Asn Ser
                                      70
                 65
Gly His Ser Phe Asn Val Asp Phe Asp Asp
                                         Thr Glu Asn Lys Ser
                 80
                                      85
                                                          90
Val Leu Arg Gly Gly Pro Leu Thr Gly Ser
                                        Tyr Arg Leu Arg Gln
                 95
                                     100
                                                          105
Val His Leu His Trp Gly Ser Ala Asp Asp His Gly Ser Glu His
                110
                                     115
Ile Val Asp Gly Val Ser Tyr Ala Ala Glu Leu His Val Val His
                125
                                     130
                                                          135
Trp Asn Ser Asp Lys Tyr Pro Ser Phe Val Glu Ala Ala His Glu
                140
                                     145
                                                          150
Pro Asp Gly Leu Ala Val Leu Gly Val Phe Leu Gln Ile Gly Glu
                155
                                     160
                                                          165
Pro Asn Ser Gln Leu Glh Lys Ile Thr Asp Thr Leu Asp Ser Ile
                                     175
                                                          180
                170
Lys Glu Lys Gly Lys Gln Thr Arg Phe Thr Asn Phe Asp Leu Leu
                                     190
                                                          195
                185
Ser Leu Leu Pro Pro Ser Trp Asp Tyr Trp Thr Tyr Pro Gly Ser
                200
                                     205
                                                          210
Leu Thr Val Pro Pro Leu Leu Glu Ser Val Thr Trp Ile Val Leu
                215
                                     220
                                                          225
Lys Gln Pro Ile Asn Ile Ser Ser Gln Gln Leu Ala Lys Phe Arg
                230
                                     235
                                                          240
Ser Leu Leu Cys Thr Ala Glu Gly Glu Ala Ala Phe Leu Val
                                     250
                                                          255
                245
His His Leu Ala Thr Ile Pro Cys Gly Ser Ser Ala Thr Cys
                                     265
                260
Ser Gly His Cys
<210> 216
<211> 182
<212> PRT
<213> Homo sapiens
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<400> 216
Lys Leu Pro Leu Pro Pro Gly Ala Phe Ser Gly Leu Trp Lys Asn
                                      10
                                                           15
Gln Glu Ala Phe Lys His Leu Tyr Phe Glu Lys Phe Pro Gly
                                      25
                 20
Tyr Asp Thr Met Asp Ala Gly Tyr Met Asp Glu Glu Gly Tyr Leu
                                      40
Tyr Val Met Ser Arg Val Asp Asp Val Ile Asn Val Ala Gly His
                                      55
                                                           60
Arg Ile Ser Ala Gly Ala Ile Glu Glu Ser Ile Leu Ser His Gly
                 65
                                      70
                                                           75
Thr Val Ala Asp Cys Ala Val Val Gly Lys Glu Asp Pro Leu Lys
                 80
                                      85
                                                           90
Gly His Val Pro Leu Ala Leu Cys Val Leu Arg Lys Asp Ile Asn
                                                          105
                 95
                                     100
Ala Thr Glu Glu Gln Val Leu Glu Glu Ile Val Lys His Val Arg
                                     115
                                                          120
                110
Gln Asn Ile Gly Pro Val Ala Ala Phe Arg Asn Ala Val Phe Val
                                     130
                                                          135
                125
Lys Gln Leu Pro Lys Thr Arg Ser Gly Lys Ile Pro Arg Ser Ala
                140
                                     145
                                                          150
Leu Ser Ala Ile Val Asn Gly Lys Pro Tyr Lys Ile Thr Ser Thr
                155
                                     160
```

Ile Glu Asp Pro Ser Ile Phe Gly His Val Glu Glu Met Leu Lys

170 175 180

Gln Ala

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<210> 217
<211> 359
<212> PRT
<213> Homo sapiens
<221> misc_feature
<223> Incyte ID No: LI:332176.1.orf2:2000FEB01
<400> 217
Leu Ile Lys Arg Ser Lys Gly Lys Glu Arg Pro Phe Val Leu Thr
                                                           15
Arg Ser Phe Phe Ala Gly Ser Gln Lys Tyr Gly Ala Val Trp Thr
                 20
                                      25
                                                           30
Gly Asp Asn Thr Ala Glu Trp Ser Asn Leu Lys Ile Ser Ile Pro
                 35
                                      40
Met Leu Leu Thr Leu Ser Ile Thr Gly Ile Ser Phe Cys Gly Ala
                                      55
                 50
Asp Ile Gly Gly Phe Ile Gly Asn Pro Glu Thr Glu Leu Leu Val
                                      70
                 65
Arg Trp Tyr Gln Ala Gly Ala Tyr Gln Pro Phe Phe Arg Gly His
                 80
                                      85
Ala Thr Met Asn Thr Lys Arg Arg Glu Pro Trp Leu Phe Gly Glu
                 95
                                     100
                                                          105
Glu His Thr Arg Leu Ile Arg Glu Ala Ile Arg Glu Arg Tyr Gly
                110
                                     115
                                                          120
Leu Leu Pro Tyr Trp Tyr Ser Leu Phe Tyr His Ala His Val Ala
                                     130
                                                          135
                125
Ser Gln Pro Val Met Arg Pro Leu Trp Val Glu Phe Pro Asp Glu
                140
                                     145
Leu Lys Thr Phe Asp Met Glu Asp Glu Tyr Met Leu Gly Ser Ala
                                     160
                                                          165
                155
Leu Trp Val His Pro Val Thr Glu Pro Lys Ala Thr Thr Val Asp
                170
                                     175
                                                          180
Val Phe Leu Pro Gly Ser Asn Glu Val Trp Tyr Asp Tyr Lys Thr
                185
                                     190
                                                          195
Phe Ala His Gly Glu Gly Gly Cys Thr Val Lys Ile Pro Val Ala
                                     205
                                                          210
                200
Leu Asp Thr Ile Pro Val Phe Gln Arg Gly Gly Ser Val Ile Pro
                                     220
                                                          225
                215
Ile Lys Thr Thr Val Gly Lys Ser Thr Gly Trp Met Thr Glu Ser
                 230
                                     235
                                                          240
Ser Tyr Gly Leu Arg Val Ala Leu Ser Thr Lys Gly Ser Ser Val
                245
                                     250
                                                          255
Gly Glu Leu Tyr Leu Asp Asp Gly His Ser Phe Gln Tyr Leu His
                                                          270
                 260
                                     265
Gln Lys Gln Phe Leu His Arg Lys Phe Ser Phe Cys Ser Ser Val
                                                          285
                 275
                                     280
Leu Ile Asn Ser Phe Ala Asp Gln Arg Gly His Tyr Pro Ser Lys
                                     295
                                                          300
                 290
Cys Val Val Glu Lys Ile Leu Val Leu Gly Phe Arg Lys Glu Pro
                 305
                                     310
                                                          315
Ser Ser Val Thr Thr His Ser Ser Asp Gly Lys Asp Gln Pro Val
                 320
                                     325
Ala Phe Thr Tyr Cys Ala Lys Thr Ser Ile Leu Ser Leu Glu Lys
                 335
                                     340
Leu Ser Leu Asn Ile Ala Thr Asp Trp Glu Val Arg Ile Ile
                 350
                                     355
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<sup>&</sup>lt;210> 218

<sup>&</sup>lt;211> 110

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> Homo sapiens

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<220>
<221> misc_feature
<223> Incyte ID No: LI:403248.2.orf2:2000FEB01
<400> 218
Ser Pro Phe Ile Ser Leu Pro Cys Ser Ala Leu Leu Lys Pro Ser
 1
                                      10
                                                           15
Thr Glu Gln Pro Leu Tyr Ser Ser Ser Leu Trp Gly Pro Ala Val
                 2.0
                                      25
                                                           30
Asp Gly Cys Asp Cys
                    Val Ala Glu Gly Leu Trp Leu Pro Gln Leu
                                      40
                                                           45
His Val Gly Asp Trp Leu Val Phe Asp Asn Met Gly Ala Tyr Thr
                 50
                                      55
                                                           60
Val Gly Met Gly Ser Pro Phe Trp Gly Thr Gln Ala Cys His Ile
                 65
                                      70
Thr Tyr Ala Met Ser Arg Val Ala Trp Glu Ala Leu Arg Arg
                                                         Gln
                 80
                                      85
Leu Met Ala Ala Glu Gln Glu Asp Asp Val Glu Gly Val Cys Lys
                 95
                                     100
                                                          105
Pro Leu Ser Cys Gly
<210> 219
<211> 549
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte ID No: LG:220992.1.orf3:2000MAY19
<400> 219
Arg Pro Val Thr Ser Phe Ser Pro Leu Pro Gly Ser Cys Gly Gly
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Arg Leu Gly Thr Arg Thr Met Leu Gly Arg Ser Leu Arg Glu Val
                 20
                                                           30
Ser Ala Ala Leu Lys Gln Gly Gln Ile Thr Pro Thr Glu Leu Cys
                 35
                                      40
                                                           45
Gln Lys Cys Leu Ser Leu Ile Lys Lys Thr Lys Phe Leu Asn Ala
                 50
                                      55
                                                           60
Tyr Ile Thr Val Ser Glu Glu Val Ala Leu Lys Gln Ala Glu Glu
                 65
                                      70
                                                           75
Ser Glu Lys Arg Tyr Lys Asn Gly Gln Ser Leu Gly Asp
                                                     Leu Asp
                 80
                                                           90
                                      85
Gly Ile Pro Ile Ala Val Lys Asp Asn Phe Ser Thr Ser Gly Ile
                                                          105
                 95
                                     100
Glu Thr Thr Cys Ala Ser Asn Met Leu Lys Gly Tyr Ile Pro Pro
                110
                                     115
                                                          120
Tyr Asn Ala Thr Val Val Gln Lys Leu Leu Asp Gln Gly Ala Leu
                                                          135
                125
                                     130
Leu Met Gly Lys Thr Asn Leu Asp Glu Phe Ala Met Gly Ser Gly
                                                          150
                                     145
                140
Ser Thr Asp Gly Val Phe Gly Pro Val Lys Asn Pro Trp Ser
                                                          Tyr
                155
                                     160
                                                          165
Ser Lys Gln Tyr Arg Glu Lys Arg Lys Gln Asn Pro His Ser Glu
                                     175
                170
                                                          180
Asn Glu Asp Ser Asp Trp Leu Ile Thr Gly Gly Ser Ser Gly Gly
                185
                                     190
                                                          195
Ser Ala Ala Ala Val Ser Ala Phe Thr Cys Tyr Ala Ala Leu Gly
                200
                                     205
                                                          210
Ser Asp Thr Gly Gly Ser Thr Arg Asn Pro Ala Ala His Cys Gly
                215
                                     220
                                                          225
Leu Val Gly Phe Lys Pro Ser Tyr Gly Leu Val Ser Arg His Gly
                                                          240
                                     235
                230
Leu Ile Pro Leu Val Asn Ser Met Asp Val Pro Gly Ile Leu Thr
                                     250
                                                          255
                245
```

Arg Cys Val Asp Asp Ala Ala Ile Val Leu Gly Ala Leu Ala Gly

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260
                                                          270
                                     265
Pro Asp Pro Arg Asp Ser Thr Thr Val His Glu Pro Ile Asn Lys
                 275
                                     280
                                                          285
Pro Phe Met Leu Pro Ser Leu Ala Asp Val Ser Lys Leu Cys Ile
                 290
                                      295
                                                          300
Gly Ile Pro Lys Glu Tyr Leu Val Pro Glu Leu Ser Ser Glu Val
                 305
                                      310
Gln Ser Leu Trp Ser Lys Ala Ala Asp Leu Phe Glu Ser Glu Gly
                 320
                                      325
                                                           330
Ala Lys Val Ile Glu Val Ser Leu Pro His Thr Ser Tyr Ser Ile
                 335
                                      340
                                                          345
Val Cys Tyr His Val Leu Cys Thr Ser Glu Val Ala Ser Asn Met
                                                          360
                 350
                                      355
Ala Arg Phe Asp Gly Leu Gln Tyr Gly His Arg Cys Asp Ile Asp
                 365
                                      370
                                                          375
Val Ser Thr Glu Ala Met Tyr Ala Ala Thr Arg Arg Glu Gly Phe
                                      385
                                                          390
                 380
Asn Asp Val Val Arg Gly Arg Ile Leu Ser Gly Asn Phe Phe Leu
                 395
                                      400
Leu Lys Glu Asn Tyr Glu Asn Tyr Phe Val Lys Ala Gln Lys Val
                                      415
                                                          420
                 410
Arg Arg Leu Ile Ala Asn Asp Phe Val Asn Ala Phe Asn Ser Gly
                 425
                                      430
                                                           435
Val Asp Val Leu Leu Thr Pro Thr Thr Leu Ser Glu Ala Val Pro
                 440
                                      445
                                                          450
 Tyr Leu Glu Phe Ile Lys Glu Asp Asn Arg Thr Arg Ser Ala Gln
                                                           465
                 455
                                      460
Asp Asp Ile Phe Thr Gln Ala Val Asn Met Ala Gly Leu Pro Ala
                 470
                                      475
Val Ser Ile Pro Val Ala Leu Ser Asn Gln Gly Leu Pro Ile Gly
                                      490
                                                           495
                 485
Leu Gln Phe Ile Gly Arg Ala Phe Cys Asp Gln Gln Leu Leu Thr
                 500
                                      505
                                                           510
Val Ala Lys Trp Phe Glu Lys Gln Val Gln Phe Pro Val Ile Gln
                 515
                                      520
                                                          525
Leu Gln Glu Leu Met Asp Asp Cys Ser Ala Val Leu Glu Asn Glu
                                      535
                                                           540
                 530
Lys Leu Ala Ser Val Ser Leu Lys Gln
                 545
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 <211> 264
 <212> PRT
 <213> Homo sapiens
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 Arg Thr Pro Ala Ala Arg Arg Pro Ala Leu Arg Phe Gly Pro Pro
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 Pro Pro Pro Thr Pro Leu Thr Leu Gly Thr Tyr Phe Gly Cys Leu
                                       25
                                                            30
                  20
 Arg Cys Pro Pro Ala Glu Thr Gln Leu Leu Arg Arg Pro Ala Val
                                       40
                  35
 Phe Val Gly Ser Ala Ala Ser Gly Ile Arg Ser Gly Leu Trp Ser
                                                            60
 Ala Ser Ser Gly His Trp Cys Ala Pro Ala Ala Gly Arg Ala His
 Ala Pro Val Pro Arg Leu Val Arg Gly Leu Gly Ala Ala Ser Thr
                  80
                                       85
                                                            90
 Ala Ala Pro Gln Asp Ala Gln Thr Gly Pro Gln Pro Met Pro Arg
                                                           105
                  95
                                      100
 Ala Asp Cys Ile Met Arg His Leu Pro Tyr Phe Cys Arg Gly Gln
                                      115
                                                           120
                 110
 Val Val Arg Gly Phe Gly Arg Gly Ser Lys Gln Leu Gly Ile Pro
```

```
125
                                     130
Thr Ala Asn Phe Pro Glu Gln Val Val Asp Asn Leu Pro Ala Asp
                140
                                     145
                                                          150
Ile Ser Thr Gly Ile Tyr Tyr Gly Trp Ala Ser Val Gly Ser Gly
                155
                                     160
                                                          165
Asp Val His Lys Met Val Val Ser Ile Gly Trp Asn Pro Tyr Tyr
                                     175
                                                          180
                170
Lys Asn Thr Lys Lys Ser Met Glu Thr His Ile Met His Thr Phe
                185
                                     190
                                                          195
Lys Glu Asp Phe Tyr Gly Glu Ile Leu Asn Val Ala Ile Val Gly
                200
                                     205
                                                          210
Tyr Leu Arg Pro Glu Lys Asn Phe Asp Ser Leu Glu Ser Leu Ile
                215
                                     220
                                                          225
Ser Ala Ile Gln Gly Asp Ile Glu Glu Ala Lys Lys Arg Leu Glu
                230
                                     235
                                                          240
Leu Pro Glu His Leu Lys Ile Lys Glu Asp Asn Phe Phe Gln Val
                245
                                     250
                                                          255
Ser Lys Ser Lys Ile Met Asn Gly His
                260
<210> 221
<211> 701
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte ID No: LI:350754.4.orf3:2000MAY01
<400> 221
Glu Glu Ala Glu Glu Gly Arg Asn Met Ala Ala Leu Gly Val Gln
                                      10
Ser Ile Asn Trp Gln Lys Ala Phe Asn Arg Gln Ala His His Thr
                                                           30
                 20
                                      25
Asp Lys Phe Ser Ser Gln Glu Leu Ile Leu Arg Arg Gly Gln Asn
                                                           45
                 35
                                      40
Phe Gln Val Leu Met Ile Met Asn Lys Gly Leu Gly Ser Asn Glu
                                      55
                 50
Arg Leu Glu Phe Ile Asp Thr Thr Gly Pro Tyr Pro Ser Glu Ser
                                      70
                                                           75
                 65
Ala Met Thr Lys Ala Val Phe Pro Leu Ser Asn Gly Ser Ser Gly
                                                           90
                                      85
                 80
Gly Trp Ser Ala Val Leu Gln Ala Ser Asn Gly Asn Thr Leu Thr
                 95
                                     100
                                                          105
Ile Ser Ile Ser Ser Pro Ala Ser Ala Pro Ile Gly Arg Tyr
                                                          Thr
                                                          120
                110
                                     115
Met Ala Leu Gln Ile Phe Ser Gln Gly Gly Ile Ser Ser Val Lys
                                     130
                                                          135
                125
Leu Gly Thr Phe Ile Leu Leu Phe Asn Pro Trp Leu Asn Val Asp
                                                          150
                140
                                     145
Ser Val Phe Met Gly Asn His Ala Glu Arg Glu Glu Tyr Val Gln
                 155
                                     160
                                                          165
Glu Asp Ala Gly Ile Ile Phe Val Gly Ser Thr Asn Arg Ile Gly
                170
                                     175
                                                          180
Met Ile Gly Trp Asn Phe Gly Gln Phe Glu Glu Asp Ile Leu Ser
                                                          195
                185
                                     190
Ile Cys Leu Ser Ile Leu Asp Arg Ser Leu Asn Phe Arg Arg Asp
                200
                                     205
                                                          210
Ala Ala Thr Asp Val Ala Ser Arg Asn Asp Pro Lys Tyr Val Gly
                                     220
                                                          225
                215
Arg Val Leu Ser Ala Met Ile Asn Ser Asn Asp Asp Asn Gly Val
                230
                                     235
                                                          240
Leu Ala Gly Asn Trp Ser Gly Thr Tyr Thr Gly Gly Arg Asp Pro
                245
                                     250
                                                          255
Arg Ser Trp Asp Gly Ser Val Glu Ile Leu Lys Asn Trp Lys Lys
                260
                                     265
                                                          270
Ser Gly Phe Ser Pro Val Arg Tyr Gly Gln Cys Trp Val Phe Ala
```

```
275
                                     280
                                                          285
Gly Thr Leu Asn Thr Ala Leu Arg Ser Leu Gly Ile Pro Ser Arg
                290
                                     295
                                                          300
Val Ile Thr Asn Phe Asn Ser Ala His Asp Thr Asp Arg Asn Leu
                305
                                     310
                                                          315
Ser Val Asp Val Tyr Tyr Asp Pro Met Gly Asn Pro Leu Asp Lys
                320
                                     325
                                                          330
Gly Ser Asp Ser Val Trp Asn Phe His Val Trp Asn Glu Gly Trp
                335
                                     340
                                                           34<del>5</del>
Phe Val Arg Ser Asp Leu Gly Pro Pro Tyr Gly Gly Trp Gln Val
                350
                                     355
                                                          360
Leu Asp Ala Thr Pro Gln Glu Arg Ser Gln Gly Val Phe Gln Cys
                365
                                     370
                                                          375
Gly Pro Ala Ser Val Ile Gly Val Arg Glu∜Gly Asp Val Gln Leu
                380
                                     385
                                                          390
Asn Phe Asp Met Pro Phe Ile Phe Ala Glu Val Asn Ala Asp Arg
                395
                                     400
                                                          405
Ile Thr Trp Leu Tyr Asp Asn Thr Thr Gly Lys Gln Trp Lys Asn
                                     415
                                                          420
                410
Ser Val Asn Ser His Thr Ile Gly Arg Tyr Ile Ser Thr Lys Ala
                425
                                     430
                                                           435
Val Gly Ser Asn Ala Arg Met Asp Val Thr Asp Lys Tyr Lys Tyr
                440
                                      445
                                                           450
Pro Glu Gly Ser Asp Gln Glu Arg Gln Val Phe Gln Lys Ala Leu
                455
                                     460
                                                           465
Gly Lys Leu Lys Pro Asn Thr Pro Phe Ala Ala Thr Ser Ser Met
                470
                                      475
                                                           480
Gly Leu Glu Thr Glu Glu Gln Glu Pro Ser Ile Ile Gly Lys Leu
                                     490
                                                           495
                485
Lys Val Ala Gly Met Leu Ala Val Gly Lys Glu Val Asn Leu Val
                500
                                      505
                                                           510
Leu Leu Lys Asn Leu Ser Arg Asp Thr Lys Thr Val Thr Val
                515
                                      520
                                                           525
Asn Met Thr Ala Trp Thr Ile Ile Tyr Asn Gly Thr Leu Val His
                530
                                      535
                                                           540
Glu Val Trp Lys Asp Ser Ala Thr Met Ser Leu Asp Pro Glu Glu
                                      550
                                                           555
                545
Glu Ala Glu His Pro Ile Lys Ile Ser Tyr Ala Gln Tyr Glu Arg
                                      565
                                                           570
                560
Tyr Leu Lys Ser Asp Asn Met Ile Arg Ile Thr Ala Val Cys
                                                          Lvs
                575
                                      580
Val Pro Asp Glu Ser Glu Val Val Glu Arg Asp Ile Ile Leu
                590
                                      595
                                                           600
Asp Asn Pro Thr Leu Thr Leu Glu Val Leu Asn Glu Ala Arg Val
                605
                                      610
                                                           615
Arg Lys Pro Val Asn Val Gln Met Leu Phe Ser Asn Pro Leu Asp
                 620
                                      625
                                                           630
Glu Pro Val Arg Asp Cys Val Leu Met Val Glu Gly Ser Gly Leu
                                      640
                                                           645
                635
Leu Leu Gly Asn Leu Lys Ile Asp Val Pro Thr Leu Gly Pro Lys
                650
                                      655
                                                           660
Glu Arg Ser Arg Val Arg Phe Asp Ile Leu Pro Ser Arg Ser Gly
                 665
                                      670
                                                           675
Thr Lys Gln Leu Leu Ala Asp Phe Ser Cys Asn Lys Phe Pro Ala
                680
                                      685
Ile Lys Ala Met Leu Ser Ile Asp Val Ala Glu
                 695
                                      700
<210> 222
<211> 150
<212> PRT
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<sup>&</sup>lt;213> Homo sapiens

<sup>&</sup>lt;220>

<sup>&</sup>lt;221> misc\_feature

<sup>&</sup>lt;223> Incyte ID No: LI:255828.29.orf2:2000MAY01

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<400> 222
Cys Thr Ile Gly Pro Ala Ser Arg Ser Val Glu Thr Leu Lys Glu
                                      10
Met Ile Lys Ser Gly Met Asn Val Ala Arg Leu Asn Phe Ser His
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                                                           30
                                      25
Gly Thr His Glu Tyr
                    His Ala Glu Thr Ile Lys Asn Val Arg Thr
                                      40
                                                           45
Ala Thr Glu Ser Phe Ala Ser Asp Pro Ile Leu Tyr Arg Pro Val
                 50
                                      55
                                                           60
Ala Val Ala Leu Asp Thr Lys Gly Pro Glu Ile Arg Thr Gly Leu
Ile Lys Gly Ser Gly Thr Ala Glu Val Glu Leu Lys Lys Ala Ala
                 80
                                      85
                                                           90
Thr Leu Lys Ile Thr Leu Asp Asn Ala Tyr Met Glu Lys Cys Asp
                 95
                                     100
                                                          105
Glu Asn Ile Leu Trp Leu Asp Tyr Lys Asn Ile Cys Lys Val Val
                110
                                     115
                                                          120
Glu Val Ser Arg Leu His His Ala Val Trp Arg Asn Ser Gln Arg
                125
                                     130
                                                          135
Gly Leu Ser Ser Gly Gly Cys Ala His Ala Ala Pro Asp Ser Ser
                140
                                     145
                                                          150
<210> 223
<211> 234
<212> PRT
<213> Homo sapiens
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Ala Ala Gly Ser Leu Phe Pro Gly Leu Leu Ile Phe Ser Met Ile
                                      10
Leu Phe Ile Phe Leu Leu Gly Tyr Ala Trp Phe Ser Ser His Thr
                 20
                                      25
                                                           30
Ser Pro Leu Tyr Trp Asp Cys Leu Leu Met Arg Gly His Glu Ile
                                      40
                                                           45
Thr Glu Gln Pro Met Lys Ala Glu Arg Ala Gly Ser Ile Met Val
                                                           60
                 50
                                      55
Lys Glu Ala Ile Ser Phe Leu Glu Arg His Ser Lys Glu Thr Phe
                                      70
                                                           75
                 65
Leu Leu Phe Phe Ser Phe Leu His Val His Thr Pro Leu Pro Thr
                 80
                                      85
                                                           90
Thr Asp Asp Phe Thr Gly Thr Ser Lys His Gly Leu Tyr Gly Asp
                 95
                                     100
                                                          105
Asn Val Asp Glu Met Asp Ser Met Val Gly Lys Ile Leu Asp Ala
                110
                                     115
                                                          120
Ile Asp Asp Phe Gly Leu Arg Asn Asn Thr Leu Val Tyr Phe Thr
                125
                                     130
                                                          135
Ser Asp His Gly Gly His Leu Glu Ala Arg Arg Gly His Ala Gln
                140
                                                          150
                                     145
Leu Gly Gly Trp Asn Gly Ile Tyr Lys Gly Gly Lys Gly Met Gly
                155
                                     160
                                                          165
Gly Trp Glu Gly Gly Ile Arg Val Pro Gly Ile Val Arg Trp Pro
                170
                                     175
                                                          180
Gly Lys Val Pro Ala Gly Arg Leu Ile Lys Glu Pro Thr Ser Leu
                                     190
                                                          195
                 185
Met Asp Ile Leu Pro Thr Val Ala Ser Val Ser Gly Gly Ser Leu
                 200
                                     205
                                                          210
Pro Gln Asp Arg Val Ile Asp Gly Arg Asp Leu Met Pro Leu Leu
                 215
                                     220
                                                          225
Ala Gly Gln Arg Gln Ala Leu Gly Ala
                230
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<210> 224

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<211> 86
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte ID No: LG:270916.2.orf2:2000FEB18
<400> 224
Lys Phe Ser Ser Asn Gly Leu Trp Pro Ser Thr Trp Leu Leu Thr
                                      10
                                                          15
Thr Arg Thr Leu Pro Met Ile Ser Arg Cys Ser Pro Met His Leu
                 20
                                      25
                                                          30
Leu Thr Ile Ser Ser Ala Phe Cys Leu Leu Cys Pro Pro Pro Arg
                 35
                                      40
                                                          45
Met Pro Phe Gln Lys Cys Leu Leu Ser Arg Tyr Arg Ser Arg
                 50
                                      55
                                                          60
Gly Val Leu Val Ala Val Ile Trp Gly Thr Thr Glu Ala Ser Gly
                 65
                                      70
Ile Ser Gly Leu Ile Thr Ala Leu Trp Glu Phe
                 80
                                      85
<210> 225
<211> 173
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte ID No: LG:999414.3.orf2:2000FEB18
<400> 225
Ile Leu Thr Ser Val Ser Ser Phe Trp Cys Pro Phe Leu
 1
                  5
                                      10
                                                           15
Ser Leu Leu Asp Ser Gln Leu His Ser Trp Ile Val Leu Gln Leu
                 20
                                      25
                                                           30
Thr Ile Ile Lys Asn Val Glu Ile Ser Asn Leu Val Cys Asp Pro
                                      40
                                                          45
                 35
Ser Gln Leu Leu Asn Leu Ala Cys Ser Asp
                                        Ser Val Ile Asp Ser
                 50
                                      55
                                                           60
Ile Phe Ile Tyr Leu Asp Ser Thr Ile Phe Gly Phe Leu Pro Ile
                                                           75
                 65
                                      70
Ser Gly Ile Leu Leu Ser Tyr Tyr Lys Ile Val Pro Ser Ile Leu
                 80
                                      85
                                                           90
Arg Ile Ser Ser Ser Asp Gly Lys Tyr Lys Ala Phe Ser Thr Cys
                 95
                                     100
                                                          105
Arg Ser His Leu Ala Val Val Cys Leu Phe Tyr Gly Thr Gly Ile
                110
                                     115
                                                          120
Gly Val Tyr Leu Thr Ser Ala Val Ala Pro Ala Pro Arg Ser
                                                         Gly
                                                          135
                125
                                     130
Val Val Val Ser Val Met Tyr Thr Val Val Thr Pro Met Leu Asn
                140
                                     145
Pro Phe Ile Tyr Cys Leu Arg Lys Gln Gly His Ser Lys Arg Leu
                155
Trp Arg Cys Ala Ala Glu Gln Ser
                170
<210> 226
<211> 68
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte ID No: LG:429446.1.orf2:2000FEB18
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<400> 226

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His Leu Val Ala Thr Val Arg Gly Phe Ser Lys Val Phe Val Ser
Ser Arg Ile Lys Thr Val Lys Leu Gln Ile Val Leu Gln Met Glu
                                                           30
                 20
                                      25
Pro Gln Met Gln Ser Met Thr Lys Ile Tyr His Arg Pro Leu Asp
                 35
                                      40
Arg Pro Ala Ser Pro Cys Ser Asp Val Asp Asp Ile Glu Gly Ala
                 50
Pro Pro Lys Glu Ile Ser Thr Ala
                 65
<210> 227
<211> 70
<212> PRT
<213> Homo sapiens
<221> misc_feature
<223> Incyte ID No: LI:057229.1.orf1:2000FEB01
<400> 227
Gln Pro Ser Leu Pro Glu Phe Ser His Phe Gln Lys Thr Val Leu
                                      10
                                                           15
Leu Glu Ser Lys Ile Ala Arg Gln Phe Ile Leu Phe Tyr Phe Ile
                 20
                                      25
                                                           30
Leu His Ile Phe Leu Arg Gln Ser Leu Ala Leu Phe Pro Arg Leu
                 35
                                      40
                                                           45
Glu Cys Gly Gly Ala Val Leu Ala His Cys Asn Leu Cys Leu Leu
                 50
Gly Ser Ser Asp Ser Pro Ala Ser Ala Ser
                 65
<210> 228
<211> 117
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte ID No: LI:351965.1.orf2:2000FEB01
<400> 228
Pro Thr Thr Ser Asn Arg Ala Ile Thr Leu Thr Ala Arg Pro Lys
                                      10
Ile Pro Phe Leu Arg Ile Arg Glu Ala Lys Asn Pro Arg Ser Glu
                                      25
Asn Met Arg Leu Ala Thr Ile Leu Glu Val Ala Cys Arg His Phe
                 35
                                      40
                                                           45
Gly Ser Gly Leu Pro Pro Ser Trp Glu Leu Trp Glu Gln Gly Pro
                                                           60
                 50
                                      55
Pro Gly Asn Ser Ser Arg Tyr Ile Glu Phe Leu Asn Lys His Thr
                                      70
                 65
Tyr Ile Lys Gly Thr Leu Arg Val Tyr Thr Lys Lys Phe Cys Met
                                      85
                                                           90
                 80
Leu Val Ile Lys Ser Phe Glu Ser Lys Ser Cys Val Cys Val Tyr
                 95
                                     100
Asp Phe Asp Ser Lys Ser Ser Val Asn Val Thr Val
<210> 229
<211> 294
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte ID No: LG:068682.1.orf2:2000FEB18
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<400> 229
Gln Arg Pro Met Ser Gly Ser Gly His Met Gly Val Arg Gly Cys
                                      10
Arg Cys Gln Ala Pro Trp Leu Arg Pro Glu Ile Gly Val Arg Pro
                                                           30
                 20
                                      25
Pro Pro Arg Ser Gln Ala Ala Ser Cys Pro Pro Cys Ala Leu Gly
                                      40
                                                           45
                 35
Ala Thr Met Ser Gly Asp Lys Leu Leu Ser Glu Leu Gly Tyr Lys
                 50
                                      55
Leu Gly Arg Thr Ile Gly Glu Gly Ser Tyr Ser Lys Val Lys Val
                 65
Ala Thr Ser Lys Lys Tyr Lys Gly Thr Val Ala Ile Lys Val Val
                                      85
                                                           90
                 80
Asp Arg Arg Ala Pro Pro Asp Phe Val Asn Lys Phe Leu Pro
                 95
                                     100
                                                          105
Arg Glu Leu Ser Ile Leu Arg Gly Val Arg His Pro His Ile Val
                110
                                     115
                                                          120
His Val Phe Glu Phe Ile Glu Val Cys Asn Gly Lys Leu Tyr Ile
                                     130
                                                          135
                125
Val Met Glu Ala Ala Ala Thr Asp Leu Leu Gln Ala Val Gln Arg
                                                          150
                140
                                     145
Asn Gly Arg Ile Pro Gly Val Gln Ala Arg Asp Leu Phe Ala Gln
                155
                                     160
                                                          165
Ile Ala Gly Ala Val Arg Tyr Leu His Asp His His Leu Val His
                170
                                     175
                                                          180
Arg Asp Leu Lys Cys Glu Asn Val Leu Leu Ser Pro Asp Glu Arg
                185
                                     190
                                                          195
Arg Val Lys Leu Thr Asp Phe Gly Phe Gly Arg Gln Ala His Gly
                                     205
                                                          210
                200
Tyr Pro Asp Leu Ser Thr Thr Tyr Cys Gly Ser Ala Ala Tyr Ala
                215
                                     220
                                                          225
Ser Pro Glu Val Leu Leu Gly Ile Pro Tyr Asp Pro Lys Lys Tyr
                230
                                     235
                                                          240
Asp Val Trp Ser Met Gly Val Val Leu Tyr Val Met Val Thr Gly
                245
                                     250
                                                          255
Cys Met Pro Phe Asp Asp Ser Asp Ile Ala Gly Leu Pro Arg Arg
                260
                                     265
                                                          270
Gln Lys Arg Gly Val Leu Tyr Pro Glu Gly Leu Glu Leu Ser Glu
                275
                                     280
                                                          285
Arg Cys Lys Ala Leu Ile Ala Glu Leu
                290
<210> 230
<211> 326
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte ID No: LG:242665.1.orf1:2000FEB18
His His Leu Ile Ser Leu Tyr Phe Thr Asp Phe Pro Ile Ser Phe
  1
                                      10
                                                           15
                                         Thr Gly Pro Ala Pro
Phe Met Phe Tyr Ala Asn Phe Ser Arg Arg
                                                           30
                  20
                                      25
Pro Leu Arg Thr Thr Pro Arg Ala Trp Leu Arg Arg Glu Cys Gly
                                                           45
                                      40
                  35
Ala Ser Thr Met Ser Ala Pro Gly Ser Pro Asp Gln Ala Tyr Asp
                                      55
                  50
Phe Leu Leu Lys Phe Leu Leu Val Gly Asp Arg Asp Val Gly Lys
                                                           75
                                      70
                  65
Ser Glu Ile Leu Glu Ser Leu Gln Asp Gly Ala Ala Glu Ser
                                                         Pro
                                      85
                                                           90
                  80
Tyr Ser His Leu Gly Gly Ile Asp Tyr Lys Thr Thr Thr Ile Leu
                  95
                                     100
Leu Asp Gly Gln Arg Val Lys Leu Lys Leu Trp Asp Thr Ser Gly
```

```
110
                                     115
                                                          120
Gln Gly Arg Phe Cys Thr Ile Phe Arg Ser Tyr Ser Arg Gly Ala
                125
                                     130
                                                          135
Gln Gly Val Ile Leu Val Tyr Asp Ile Ala Asn Arg Trp Ser Phe
                140
                                     145
Glu Gly Met Asp Arg Trp Ile Lys Lys Ile Glu Glu His Ala Pro
                155
                                     160
Gly Val Pro Lys Ile Leu Val Gly Asn Arg Leu His Leu Ala Phe
                170
                                     175
                                                          180
Lys Arg Gln Val Pro Arg Glu Gln Ala Gln Ala Tyr Ala Glu Arg
                185
                                     190
                                                          195
Leu Gly Val Thr Phe Phe Glu Val Ser Pro Leu Cys Asn Phe Asn
                200
                                     205
                                                          210
Ile Ile Glu Ser Phe Thr Glu Leu Ala Arg Ile Val Leu Leu Arg
                215
                                     220
                                                          225
His Arg Met Asn Trp Leu Gly Arg Pro Ser Lys Val Leu Ser Leu
                                     235
                                                          240
                230
Gln Asp Leu Cys Cys Arg Thr Ile Val Ser Cys Thr Pro Val His
                                     250
                245
Leu Val Asp Lys Leu Pro Leu Pro Ser Thr Leu Arg Ser His Leu
                260
                                     265
                                                          270
Lys Ser Phe Ser Met Ala Lys Gly Leu Asn Ala Arg Met Met Arg
                275
                                     280
                                                          285
Gly Leu Ser Tyr Ser Leu Thr Thr Ser Ser Thr His Lys Arg Ser
                290
                                     295
                                                          300
Ser Leu Cys Lys Val Lys Ile Val Cys Pro Pro Gln Ser Pro Pro
                305
                                     310
                                                          315
Lys Asn Cys Thr Arg Asn Ser Cys Lys Ile Ser
                320
<210> 231
<211> 182
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte ID No: LG:241743.1.orf1:2000FEB18
<400> 231
Lys Ser Gly Thr Pro Arg Arg Ala Leu Leu Leu Phe Leu Val Phe
                                      10
Lys Ile Arg Gly Ser Pro Val Ser His Leu Met Pro Arg Leu Lys
                                      25
                  20
Glu Ser Arg Ser His Glu Ser Leu Leu Ser Pro Ser Ser Ala Val
                                      40
                  35
Glu Ala Leu Asp Leu Ser Met Glu Glu Glu Val Val Ile Lys Pro
                  50
                                      55
                                                           60
Val His Ser Ser Ile Leu Gly Gln Asp Tyr Cys Phe Glu Val Thr
                  65
                                      70
                                                           75
Thr Ser Ser Gly Ser Lys Cys Phe Ser Cys Arg Ser Ala Ala Glu
                                                           90
                                      85
                  80
Arg Asp Lys Trp Met Glu Asn Leu Arg Arg Ala Val His Pro Asn
                                     100
                                                          105
                  95
Lys Asp Asn Ser Arg Arg Val Glu His Ile Leu Lys Leu Trp Val
                                     115
                                                          120
                 110
Ile Glu Ala Lys Asp Leu Pro Ala Lys Lys Lys Tyr Leu Cys Glu
                 125
                                     130
Leu Cys Leu Asp Asp Val Leu Tyr Ala Arg Thr Thr Gly Lys Leu
                 140
                                      145
                                                          150
Lys Thr Asp Asn Val Phe Trp Gly Glu His Phe Glu Phe His Asn
                 155
                                     160
                                                          165
Leu Pro Pro Leu Arg Thr Val Thr Val His Leu Tyr Arg Glu Thr
                 170
                                     175
                                                          180
Asp Lys
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<210> 232
<211> 358
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte ID No: LI:034212.1.orf1:2000FEB01
<221> unsure
<222> 25
<223> unknown or other
<400> 232
Asn Ser Ser Leu Thr Gln Leu Arg Arg Leu Glu Glu Leu Asp Leu
                                      10
Gly Asn Asn Glu Ile Tyr Asn Leu Pro Xaa Ser Ile Gly Ala Leu
                 20
                                      25
Leu His Leu Lys Asp Leu Trp Leu Asp Gly Asn Gln Leu Ser Glu
                                      40
                                                           45
Leu Pro Gln Glu Ile Gly Asn Leu Lys Asn Leu Leu Cys Leu Asp
                 50
                                      55
                                                           60
Val Ser Glu Asn Arg Leu Glu Arg Leu Pro Glu Glu Ile Ser Gly
                                                           75
                 65
                                      70
Leu Thr Ser Leu Thr Asp Leu Val Ile Ser Gln Asn Leu Leu Glu
                                                           90
                 80
                                      85
Thr Ile Pro Asp Gly Ile Gly Lys Leu Lys Lys Leu Ser Ile Leu
                 95
                                     100
                                                          105
Lys Val Asp Gln Asn Arg Leu Thr Gln Leu Pro Glu Ala Val Gly
                110
                                     115
                                                          120
Glu Cys Glu Ser Leu Thr Glu Leu Val Leu Thr Glu Asn Gln Leu
                125
                                     130
Leu Thr Leu Pro Lys Ser Ile Gly Lys Leu Lys Lys Leu Ser Asn
                140
                                     145
                                                          150
Leu Asn Ala Asp Arg Asn Lys Leu Val Ser Leu Pro Lys Glu Ile
                155
                                     160
                                                          165
Gly Gly Cys Cys Ser Leu Thr Val Phe Cys Val Arg Asp Asn Arg
                                     175
                                                          180
                170
Leu Thr Arg Ile Pro Ala Glu Val Ser Gln Ala Thr Glu Leu His
                                     190
                185
Val Leu Asp Val Ala Gly Asn Arg Leu Leu His Leu Pro Leu Ser
                200
                                     205
                                                          210
Leu Thr Ala Leu Lys Leu Lys Ala Leu Trp Leu Ser Asp Asn Gln
                215
                                     220
                                                          225
Ser Gln Pro Leu Leu Thr Phe Gln Thr Asp Thr Asp Tyr Thr Thr
                                                          240
                230
                                     235
Gly Glu Lys Ile Leu Thr Cys Val Leu Leu Pro Gln Leu Pro Ser
                245
                                     250
                                                          255
Glu Pro Thr Cys Gln Glu Asn Leu Pro Arg Cys Gly Ala Leu Glu
                                     265
                                                          270
                260
Asn Leu Val Asn Asp Val Ser Asp Glu Ala Trp Asn Glu Arg Ala
                275
                                     280
                                                          285
Val Asn Arg Val Ser Ala Ile Arg Phe Val Glu Asp Glu Lys Asp
                290
                                     295
                                                          300
Glu Glu Asp Asn Glu Thr Arg Thr Leu Leu Arg Arg Ala Thr Pro
                305
                                     310
                                                          315
His Pro Gly Glu Leu Lys His Met Lys Lys Thr Val Glu Asn Leu
                320
                                     325
                                                          330
Arg Asn Asp Met Asn Ala Ala Lys Gly Leu Asp Ser Asn Lys Asn
                                     340
                335
Glu Val Asn His Ala Ile Asp Arg Val Thr Thr Ser Val
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                                     355
<210> 233
<211> 194
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<sup>&</sup>lt;212> PRT

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<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte ID No: LG:344886.1.orf1:2000MAY19
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Glu Lys Met Gly Lys Gly Cys Lys Val Val Val Cys Gly Leu Leu
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Ser Val Gly Lys Thr Ala Ile Leu Glu Gln Leu Leu Tyr Gly Asn
                 20
                                      25
                                                          30
His Thr Ile Gly Met. Glu Asp Cys Glu Thr Met Glu Asp Val Tyr
                 35
                                      40
Met Ala Ser Val Glu Thr Asp Arg Gly Val Lys Glu Gln Leu Hís
                                      55
                 50
                                                           60
Leu Tyr Asp Thr Arg Gly Leu Gln Glu Gly Val Glu Leu Pro
                                                         Lys
                                      70
                                                           75
                 65
His Tyr Phe Ser Phe Ala Asp Gly Phe Val Leu Val Tyr Ser Val
                 80
                                      85
                                                           90
Asn Asn Leu Glu Ser Phe Gln Arg Val Glu Leu Leu Lys Lys Glu
                                     100
                 95
                                                         105
Ile Asp Lys Phe Lys Asp Lys Lys Glu Val Ala Ile Val Val Leu
                110
                                     115
                                                         120
Gly Asn Lys Ile Asp Leu Ser Glu Gln Arg Gln Val Asp Ala Glu
                                                          135
                                     130
                125
Val Ala Gln Gln Trp Ala Lys Ser Glu Lys Val Arg Leu Trp Glu
                                     145
                                                          150
                140
Val Thr Val Thr Asp Arg Lys Thr Leu Ile Glu Pro Phe Thr Leu
                155
                                     160
                                                          165
Leu Ala Ser Lys Leu Ser Gln Pro Gln Ser Lys Ser Ser Phe Pro
                170
                                     175
                                                          180
Leu Pro Gly Arg Lys Asn Lys Gly Asn Ser Asn Ser Glu Asn
                185
                                     190
<210> 234
<211> 222
<212> PRT
<213> Homo sapiens
<221> misc_feature
<223> Incyte ID No: LG:228930.1.orf2:2000MAY19
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Ala Gln Met Ala Gly Ala Gln Pro Gly Val His Ala Leu Gln Leu
                                      10
                                                           15
Lys Pro Val Cys Val Ser Asp Ser Leu Lys Lys Gly Thr Lys Phe
                                                           3.0
                 20
                                      25
Val Lys Trp Asp Asp Asp Ser Thr Ile Val Thr Pro Ile Ile Leu
                 35
                                      40
                                                           45
Arg Thr Asp Pro Gln Gly Phe Phe Tyr Trp Thr Asp Gln Asn
                                      55
Lys Glu Thr Glu Leu Leu Asp Leu Ser Leu Val Lys Asp Ala Arg
                                      70
                                                           75
                 65
Cys Gly Arg His Ala Lys Ala Pro Lys Asp Pro Lys Leu Arg Glu
                 80
                                      85
                                                           90
Leu Leu Asp Val Gly Asn Ile Gly Arg Leu Glu Gln Arg Met Ile
                                                          105
                 95
                                     100
Thr Val Val Tyr Gly Pro Asp Leu Val Asn Ile Ser His Leu Asn
                110
                                     115
Leu Val Ala Phe Gln Glu Glu Val Ala Lys Glu Trp Thr Asn Glu
                                     130
                                                          135
                125
Val Phe Ser Leu Ala Thr Asn Leu Leu Ala Gln Asn Met Ser Arg
                140
                                     145
                                                          150
Asp Ala Phe Leu Glu Lys Ala Tyr Thr Lys Leu Lys Leu Gln Val
                155
                                     160
```

Thr Pro Glu Gly Arg Ile Pro Leu Lys Asn Ile Tyr Arg Leu Phe

```
180
                170
                                     175
Ser Ala Asp Arg Lys Arg Val Glu Thr Ala Leu Glu Ala Cys Ser
                185
                                    . 190
Leu Pro Ser Ser Arg Val Glu Lys Ala Asn Glu Ala Ala Lys Ser
                200
                                     205
Glu Gln Ser Cys Gly Lys Ala Pro Pro Lys His Phe
                215
<210> 235
<211> 185
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte ID No: LG:338927.1.orf3:2000MAY19
<400> 235
Leu Arg Ser Thr Pro Glu Thr Gly Arg Met Lys Gly Ala Ser Glu
 1
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Glu Lys Leu Ala Ser Val Ser Asn Leu Val Thr Val Phe Glu Asn
                                                           30
                 20
                                      25
Ser Arg Thr Pro Glu Ala Ala Pro Arg Gly Gln Arg Leu Glu Asp
                                      40
                 35
Val His His Arg Pro Glu Cys Arg Pro Pro Glu Ser Pro Gly Pro
                 50
                                                           60
Arg Glu Lys Thr Asn Val Gly Glu Ala Val Gly Ser Glu Pro Arg
                 65
                                      70
                                                           75
Thr Val Ser Arg Arg Tyr Leu Asn Ser Leu Lys Asn Lys Leu Ser
                                                           90
                 80
                                      85
Ser Glu Ala Trp Arg Lys Ser Cys Gln Pro Val Thr Leu Ser Gly
                 95
                                                          105
                                     100
Ser Gly Thr Gln Glu Pro Glu Lys Lys Ile Val Gln Glu Leu Leu
                                     115
                                                          120
                110
Glu Thr Glu Gln Ala Tyr Val Ala Arg Leu His Leu Leu Asp Gln
                                      130
                                                          135
                125
Val Phe Phe Gln Glu Leu Leu Lys Thr Ala Arg Ser Ser Lys Ala
                140
                                     145
                                                          150
Phe Pro Glu Asp Val Val Arg Val Ile Phe Ser Asn Ile Ser Ser
                155
                                     160
                                                          165
Ile Tyr Gln Phe His Ser Gln Phe Phe Leu Pro Glu Leu Gln Arg
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                                     175
                Trp
Arg Leu Asp Asp
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Arg Pro Leu Glu His Gly Thr His Arg His Ile Ala Ser Leu Lys
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Thr Glu Glu Thr Arg Arg Ala Arg Pro Ala Ala Ala Gln Ala Val
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Tyr Leu Pro Val Ser Gln His Gly His Gln Asp Pro Val His Phe
                                                            45
                                       40
                 35
Ala Leu Ser Gln Arg Arg Gly Pro Ser Leu Pro Ala Ala Ala Thr
                  50
                                       55
Val Pro Pro Asp Leu Pro Ser Glu Asp Pro His Pro Gly Ala Gly
                  65
                                       70
                                                            75
Pro Pro Glu His Gly Gln Pro Arg Pro Leu Pro Asp Gly His His
                  80
                                       85
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Gln Cys Pro Gln Leu Leu Pro Ser Gln Ser Thr Arg Cys Arg Leu
                 95
                                     100
Leu Gln Leu Pro Leu Cys Ala Glu Arg Asp Leu Gly Pro Ala Ala
                110
                                     115
                                                          120
Gly Ser Arg Val Cys Ser Lys Gly Arg Val Gly Ala Ala Gly Arg
                125
                                     130
                                                          135
His Val Trp Arg Arg Gln Pro Gln Gly Leu Ser Pro Pro Gly Ala
                140
                                     145
                                                          150
Val Val His Leu Val Thr Gln Asp Arg Ala Ile Val Thr Arg Arg
                155
                                     160
                                                          165
Gly Arg His Arg Gln Pro Arg Ala Cys Gly Arg Val Leu Glu Val
                                                          180
                170
                                     175
Val Ser Ala His Arg Glu Trp Ser Arg Ser Trp Arg
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                                     190
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Ala His Ser Lys Pro Glu Lys Ile Val Asn Lys Pro Asn Lys His
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Glu Asp His Thr Gly Lys Leu Arg Pro Glu Thr Arg Glu Glu Asn
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Lys Asn His Leu Lys Asp His Gln Pro Tyr Trp His Thr Phe
                                                         Val
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Asn Asn Thr Gln Phe Pro Asp Ile Trp Glu Gln Val Lys Cys Val
Thr
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Arg Cys Gly Ala Ala Ala Ser Ala Gly Arg Glu Ser Ala Ala Gly
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Ser Glu Glu Gln Ala Gly Leu Arg Pro Ser Gln Leu Arg Gly Pro
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                                      25
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Pro Asp Pro Pro Thr Glu Thr Ala Ala Val Ser Gly Gln Ala Val
                 35
                                      40
                                                           45
Gly Ala Ala Trp Pro Ala Ala Gly Lys Met Phe Ser Val Glu Ser
                  50
                                      55
                                                           60
Leu Glu Arg Ala Glu Leu Cys Glu Ser Leu Leu Thr Trp Ile Gln
                                                           75
                                      70
                  65
Thr Phe Asn Val Gly Cys Thr Met Pro Glu Pro Val Glu Asp Leu
                                      85
                                                           90
                 80
Thr Asn Gly Val Val Met Ala Gln Val Leu Gln Lys Ile Asp Pro
                  95
                                     100
                                                          105
Ala Tyr Phe Asp Glu Asn Trp Leu Asn Arg Ile Lys Thr Glu Val
                110
                                     115
Gly Asp Asn Trp Arg Leu Lys Ile Ser Asn Leu Lys Lys Ile Leu
                125
                                     130
                                                          135
Lys Gly Ile Leu Asp Tyr Asn His Glu Ile Leu Gly Gln Gln Ile
                140
                                     145
                                                          150
Asn Asp Phe Thr Leu Pro Asp Val Asn Leu Ile Gly Glu His Ser
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155
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Asp Ala Ala Glu Leu Gly Arg Met Leu Gln Leu Ile Leu Gly Cys
                170
                                     175
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Ala Val Asn Cys Glu Gln Lys Gln Glu Tyr Ile Gln Ala Ile Met
                185
                                     190
                                                         195
Met Met Glu Glu Ser Val Gln His Val Val Met Thr Ala Ile Gln
                200
                                     205
                                                         210
Glu Leu Met Ser Lys Glu Ser Pro Val Ser Ala Gly Asn Asp Ala
                                                         225
                                     220
                215
Tyr Val Asp Leu Asp Arg Gln Leu Lys Lys Thr Thr Glu Glu Leu
                                                         240
                                     235
                230
Asn Glu Ala Leu Ser Ala Lys Glu Glu Ile Ala Gln Arg Cys His
                245
                                     250
                                                         255
Glu Leu Asp Met Gln Val Ala Ala Leu Gln Glu Glu Lys Ser Ser
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                260
                                     265
Leu Leu Ala Glu Asn Gln Val Leu Met Glu Arg Leu Asn Gln Ser
                                                          285
                275
                                     280
Asp Ser Ile Glu Asp Pro Asn Ser Pro Ala Gly Arg Arg His Leu
                290
                                     295
                                                          300
Gln Leu Gln Thr Gln Leu Glu Gln Leu Gln Glu Glu Thr Phe Arg
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                305
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Leu Glu Ala Ala Lys Asp Asp Tyr Arg Ile Arg Cys Glu Glu Leu
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Glu Lys Gly Asp Leu
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Ala His Cys Ser Arg Ala Cys Ala Ala Cys Thr Ser Pro Arg Thr
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Ala Cys Arg Thr Leu Thr Ala Ser Ser Ala Pro Ser Pro Trp Thr
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                                      40
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Ser Ser Leu Pro Thr Pro Leu Ala Gly Gly Pro Thr Ala Pro Gly
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                                      55
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Pro Pro Thr Pro Ala Arg Pro Arg Ser Ser Ala Ser Trp Thr Ala
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                                                           75
                 65
Arg Arg Gly Pro Arg Trp Ala Cys Pro Arg Pro Ala Arg Thr Ala
                                                           90
                 80
                                      85
Arg Thr Pro Arg Leu Arg Arg Gly Pro Arg Pro Arg Arg Arg Pro
                                                          105
                                     100
                 95
Arg Pro Pro Ala Gly Ser Pro Ala Arg Ser Pro Ala His Ser Leu
                                     115
                110
Gly Leu Asn Phe Gly Asp Ala Ala Arg Gln Thr Pro Arg His Gly
                                                          135
                125
                                     130
Leu Ser Ala Leu Ser Ala Pro Gly Leu Pro Gly Pro Gly Gln Pro
                                                          150
                140
                                     145
Ala Gly Pro Gly Ala Trp Ala Pro Pro Leu Asp Ser Pro Gly Thr
                                     160
                                                          165
                 155
Pro Ser Pro Asp Gly Pro Trp Cys Phe Ser Pro Glu Gly Ala Gln
                170
                                     175
                                                          180
Gly Ala Gly Gly Val Leu Phe Ala Pro Phe Gly Arg Ala Gly Ala
                                                          195
                185
                                     190
Pro Gly Pro Gly Gly Ser Asp Leu Arg Arg Glu Ala Ala
                                                          210
                                     205
                200
Arg Ala Glu Pro Arg Asp Ala Arg Thr Gly Trp Pro Glu Glu Pro
                                     220
                215
Ala Pro Glu Thr Gln Phe Lys Arg Arg Ser Cys Gln Met Glu Phe
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230
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Glu Glu Gly Met Val Glu Gly Arg Ala Arg Gly Glu Glu Leu Ala
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Ala Leu Gly Lys Gln Ala Ser Phe Ser Gly Asn Val Glu Val Ile
                                     265
                                                         270
                260
Gln Val Val Ser Asp Pro Ser Ala Ala Phe Gly Pro Ala Ala Arg
                275
                                     280
                                                          285
Ser Gln Ala Arg Asn Lys Cys Ile Leu Tyr Ile Met Gln Arg Lys
                                     295
                290
Val Asn Gly Leu Thr Gly Asn Phe Asn Pro Arg Ser Lys Ile Ile
                305
                                     310
Ser Ile Phe Tyr Leu Phe Lys Leu Phe Ile Leu Ala Met Asp Leu
                320
                                     325
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Ala Thr Val Arg Val Val Leu Glu Leu Pro Phe Leu Leu Ser Gly
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                                     340
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Ala Thr Trp Ala Phe Ile Ser Ala Pro Val Pro Val Phe Pro Asp
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Ser Phe Gly Ile Lys Ala Ser Ser Glu Ala Ser Thr Leu Glu Ala
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Met Gly Arg Lys Glu Glu Asp Asp Cys Ser Ser Trp Lys Lys Gln
                                                           45
                                      40
                 35
Thr Thr Asn Ile Arg Lys Thr Phe Ile Phe Met Glu Val Leu Gly
                 50
                                      55
                                                           60
Ser Gly Ala Phe Ser Glu Val Phe Leu Val Lys Gln Arg Leu Thr
                 65
Gly Lys Leu Phe Ala Leu Lys Cys Ile Lys Lys Ser Pro Ala Phe
                 80
                                      85
                                                           90
Arg Asp Ser Ser Leu Glu Asn Glu Ile Ala Val Leu Lys Lys Ile
                                                          105
                 95
                                     100
Lys His Glu Asn Ile Val Thr Leu Glu Asp Ile Tyr Glu Ser Thr
                110
                                     115
                                                          120
Thr His Tyr Tyr Leu Val Met Gln Leu Val Ser Gly Gly Glu Leu
                                     130
                                                          135
                125
Phe Asp Arg Ile Leu Glu Arg Gly Val Tyr Thr Glu Lys Asp Ala
                140
                                     145
Ser Leu Val Ile Gln Gln Val Leu Ser Ala Val Lys Tyr Leu His
                                     160
                                                          165
                155
Glu Asn Gly Ile Val His Arg Asp Leu Lys Pro Glu Asn Leu Leu
                170
                                     175
Tyr Leu Thr Pro Glu Glu Asn Ser Lys Ile Met Ile Thr Asp Phe
                                                          195
                185
                                     190
Gly Leu Ser Lys Met Glu Gln Asn Gly Ile Met Ser Thr Ala Cys
                                                          210
                200
                                     205
Gly Thr Pro Gly Tyr Val Ala Pro Glu Val Leu Ala Gln Lys Pro
                215
                                     220
Tyr Ser Lys Ala Val Asp Cys Trp Ser Ile Gly Val Ile Thr Tyr
                230
                                     235
Ile Leu Leu Cys Gly Tyr Pro Pro Phe Tyr Glu Glu Thr Glu Ser
                245
                                     250
                                                          255
Lys Leu Phe Glu Lys Ile Lys Glu Gly Tyr Tyr Glu Phe Glu Ser
                260
                                     265
                                                          270
Pro Phe Trp Asp Asp Ile Ser Glu Ser Ala Lys Asp Phe Ile Cys
                275
                                     280
                                                          285
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His Leu Leu Glu Lys Asp Pro Asn Glu Gly Val Thr Leu

PCT/US01/06059

30

45

60

75

90

Tle

105

120

WO 01/62927 290 295 <210> 241 <211> 133 <212> PRT <213> Homo sapiens <220> <221> misc\_feature <223> Incyte ID No: LI:338927.1.orf1:2000MAY01 <400> 241 Asp Pro Pro Arg Glu Thr Gly Arg Met Lys Gly Ala Ser Glu Glu 10 Lys Leu Ala Ser Val Ser Asn Leu Val Thr Val Phe Glu Asn Ser 20 25 Arg Thr Pro Glu Ala Ala Pro Arg Gly Gln Arg Leu Glu Asp Val 40 35 His His Arg Pro Glu Cys Arg Pro Pro Glu Ser Pro Gly Pro Arg 50 55 Glu Lys Thr Asn Val Gly Glu Ala Val Gly Ser Glu Pro Arg Thr 65 70 Val Ser Arg Arg Tyr Leu Asn Ser Leu Lys Asn Lys Leu Ser Ser 80 85 Glu Ala Trp Arg Lys Ser Leu Pro Ala Cys Asp Pro Leu Arg 95 100 Gly Asp Ala Gly Ala Arg Glu Glu Asp Arg Pro Gly Ala Ala Trp 110 115 Arg His Asp Ala Gly His Met Trp Arg Ala Ser Thr Cys 125 <210> 242 <211> 354

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160
Ile Glu Val Thr Ser Glu Arg Glu Ser Thr Ser Gly Ser Ser Val
                170
                                                          180
                                     175
Ser Ser Ser Val Ser Thr Leu Ser His Tyr Thr Glu Ser Val Gly
                                     190
                                                          195
                185
Ser Gly Glu Met Gly Ala Pro Thr Val Val Val Gln Gln Gln Gln
                200
                                     205
                                                          210
Gln Gln Gln Arg Leu Leu Gln Gln Gln Pro Ala Leu Gln Gly Val
                215
                                     220
Thr Leu Gln Gln Met Asp Phe Gly Ser Thr Gly Pro Gln Ser Ile
                230
                                     235
                                                          240
Pro Ala Val Ser Ile Pro Gln Ser Ile Ser Gln Ser Gln Ile Ser
                245
                                     250
                                                          255
Gln Val Gln Leu Gln Ser Gln Glu Leu Ser Tyr Gln Gln Lys Gln
                260
                                     265
                                                          270
Gly Leu Gln Pro Val Pro Leu Gln Ala Thr Met Ser Ala Ala Thr
                                                          285
                275
                                     280
Gly Ile Gln Pro Ser Pro Val Asn Val Val Gly Val Thr Ser Ala
                                     295
                290
                                                          300
Leu Gly Gln Gln Pro Ser Ile Ser Ser Leu Ala Gln Pro Gln Leu
                305
                                     310
                                                          315
Pro Tyr Ser Gln Ala Ala Pro Pro Val Gln Thr Pro Leu Pro Gly
                                     325
                320
                                                          330
Ala Pro Pro Pro Gln Gln Leu Gln Tyr Gly Xaa Gln Gln Pro Met
                                     340
                335
Val Ser Thr Gln Met Ala Pro Gly Met
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Leu Tyr Cys Ile Cys Lys Thr Pro Tyr Asp Glu Ser Lys Phe Tyr
Ile Gly Cys Asp Leu Cys Thr Asn Trp Tyr
                                         His Gly Glu Cys Val
                 20
                                      25
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Gly Ile Thr Glu Lys Glu Ala Lys Lys Met Asp Val Tyr Ile Cys
                                                           45
                                      40
Asn Asp Cys Lys Arg Ala Gln Glu Gly Ser Ser Glu Glu Leu Tyr
                                                           60
                 50
                                      55
Cys Ile Cys Arg Thr Pro Tyr Asp Glu Ser Gln Phe Tyr Ile Gly
                                      70.
                 65
Cys Asp Arg Cys Gln Asn Trp Tyr His Gly Arg Cys Val Gly Ile
                 80
                                      85
Leu Gln Ser Glu Ala Glu Leu Ile Asp Glu Tyr Val Cys Pro Gln
                 95
                                     100
                                                          105
Cys Gln Ser Thr Glu Asp Ala Met Thr Val Leu Thr Pro Leu Thr
                110
                                     115
                                                          120
Glu Lys Asp Tyr Glu Gly Leu Lys Arg Val Leu Arg Ser Leu Gln
                125
                                     130
                                                          135
Ala His Lys Met Ala Trp Pro Phe Leu Glu Pro Val Asp Pro Asn
                                                          150
                140
                                     145
Asp Ala Pro Asp Tyr Tyr Gly Val Ile Lys Glu Pro Met Asp Leu
                155
                                     160
                                                          165
Ala Thr Met Glu Glu Arg Val Gln Arg Arg Tyr Tyr Glu Lys Leu
                170
                                     175
Thr Glu Phe Val Ala Asp Met Thr Lys Ile Phe Asp Asn Cys Arg
                185
                                     190
                                                          195
Tyr Tyr Asn Pro Ser Asp Ser Pro Phe Tyr Gln Cys Ala Glu Val
                200
                                     205
                                                          210
Leu Glu Ser Phe Phe Val Gln Lys Leu Lys Gly Phe Lys Ala Ser
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220
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                215
Arg Ser His Asn Asn Lys Leu Gln Ser Thr Ala Ser
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Pro Pro Trp Gly Gln Arg Ser Pro Thr Pro Pro Ser Asp Thr Gly
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Gly Thr Ser Arg Pro Arg Thr Met Ile Pro Pro Gly Glu Cys Thr
                                                           30
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Tyr Ala Gly Arg Lys Arg Arg Arg Pro Leu Gln Lys Gln Arg Pro
                 35
                                      40
Ala Val Gly Ala Glu Lys Ser Asn Pro Ser Lys Arg His Arg Asp
                                      55
                                                           60
                 50
Arg Leu Asn Ala Glu Leu Asp His Leu Ala Ser Leu Leu Pro Phe
                 65
                                      70
                                                           75
Pro Pro Asp Ile Ile Ser Lys Leu Asp Lys Leu Ser Val Leu Arg
                                                           90
                 80
                                      85
Leu Ser Val Ser Tyr Leu Arg Val Lys Ser Phe Phe Gln Gly Gln
                 95
                                     100
                                                          105
Gly Leu Ala Val Ala Asp Ala Glu Asp Val Asp Asp His Thr Gly
                                     115
                                                          120
                110
Glu Arg Arg Pro Met Ser Phe Arg Arg Pro Arg Ala Leu Asp Thr
                                     130
                125
                                                          135
Gln Ala Leu Arg Arg Thr Gln Phe Gly Leu His Leu Leu Met Val
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                                     145
                                                          150
Asn Ile Ala Gly Leu Ile Ala Thr Asp Arg Leu
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His Glu Phe Gly Ala Glu Gly Pro Trp Gly Asn Met Ala Glu Gly
                 20
                                      2.5
                                                           30
Gly Ala Ser Lys Gly Gly Glu Glu Pro Gly Lys Leu Pro Glu
                                                           45
                                      40
                 35
Pro Ala Glu Glu Ser Gln Val Leu Arg Gly Thr Gly His Cys
                                      55
                 50
Lys Trp Phe Asn Val Arg Met Gly Phe Gly Phe Ile Ser Met
                                                          Ile
                                      70
                 65
Asn Arg Glu Gly Ser Pro Leu Asp Ile Pro Val Asp Val Phe Val
                                                           90
                 80
                                      85
His Gln Ser Lys Leu Phe Met Glu Gly Phe Arg Ser Leu Lys Glu
                 95
                                     100
                                                          105
Gly Glu Pro Val Glu Phe Thr Phe Lys Lys Ser Ser Lys Gly Leu
                 110
                                     115
                                                          120
Glu Ser Ile Arg Val Thr Gly Pro Gly Gly Ser Pro Cys Leu Gly
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                125
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Ser Glu Arg Arg Pro Lys Gly Lys Thr Leu Gln Lys Arg Lys Pro
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                                     145
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Lys

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<223> unknown or other
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Met Ala Ser Lys Val Thr Asp Ala Ile Val Trp Tyr Gln Lys Lys
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Ile Gly Ala Tyr Asp Gln Gln Ile Trp Glu Lys Ser Val Glu Gln
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                 20
                                      25
Arg Glu Ile Lys Gly Leu Arg Asn Lys Pro Lys Lys Thr Ala His
                                                           45
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                                      40
Val Lys Pro Asp Leu Ile Asp Val Asp Leu Val Arg Gly Ser Ala
                 50
                                      55
                                                           60
Phe Ala Lys Ala Lys Pro Glu Ser Pro Trp Thr Ser Leu Thr Arg
                  65
                                      70
                                                           75
Lys Gly Ile Val Arg Val Val Phe Phe Pro Phe Phe Arg
                                                          Trp
                  80
                                      85
                                                           90
Trp Leu Gln Val Thr Ser Lys Val Ile Phe Phe Trp Leu Leu Val
                 95
                                     100
                                                          105
Leu Tyr Leu Leu Gln Val Ala Val Ile Val Leu Phe Cys Ser Thr
                                     115
                                                          120
                 110
Ser Ser Pro His Ser Ile Pro Leu Thr Glu Val Ile Gly Pro Ile
                                     130
                 125
                                                          135
Trp Leu Met Leu Leu Gly Thr Val His Cys Gln Ile Val Ser
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                                     145
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Thr Arg Thr Pro Lys Pro Xaa Leu Ser Thr
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Leu His Ile Pro Arg Ser Pro Pro Gly Asp Arg Ala Ala Arg Thr
Gly His Pro Arg Leu Pro Val Pro Pro Pro Arg Ala Arg Thr Glu
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Pro Arg Pro Arg Gly Gln Arg Arg Leu His Ser Ser Gly Glu Met
                  35
                                      40
                                                           45
Ala Ala Gly Ser Thr Thr Leu Arg Ala Val Gly Lys Leu Gln Val
                  50
                                                           60
                                      55
Arg Leu Ala Thr Lys Thr Glu Pro Lys Lys Leu Glu Lys Tyr Leu
                                                           75
                  65
                                      70
Gln Lys Leu Ser Ala Leu Pro Met Thr Ala Asp Ile Leu Ala Glu
                  80
                                      85
Thr Gly Leu Arg Lys Thr Val Lys Arg Leu Arg Lys His Gln His
                                     100
Val Gly Asp Phe Ala Arg Asp Leu Ala Ala Arg Trp Lys Lys Leu
                                     115
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Val Leu Val Asp Arg Asn Thr Gly Pro Asp Pro Gln Asp Pro Glu
                125
                                    130
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Glu Ser Ala Ser Arg Gln Arg Tyr Gly Glu Ala Leu Gln Glu Arg
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Glu Lys Gly Trp Gly Leu Pro Arg Lys Arg
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Arg Ala His Ser Gly Val Leu Met Ser Ala Met Leu Ser His Gly
Val Leu Lys Arg Ala Ser Glu Arg Gly Ala Glu Arg His Ser Leu
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Pro Pro Ser Arg Leu Val Leu Val Pro Gly Arg Arg Ala Leu Arg
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Ser Ala Pro Gln Val Pro Gly Ser Gly Trp Arg Val Gly Thr Glu
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                                                          60
Pro Pro Val Leu His Asp Pro Ala Gly Arg Gly Arg Phe Pro Gln
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                                      70
Ser Gly Glu Val Ser Ala Ala Pro Glu Met
                                        Ser Lys Leu Ser Phe
                 80
                                     85
                                                          90
Arg Ala Arg Ala Leu Asp Ala Ser Lys Pro Leu Pro Val Phe Arg
                                    100
                 95
Cys Glu Asp Leu Pro Asp Leu His Glu Tyr Ala Ser Ile Asn Arg
                                    115
                110
Ala Val Pro Gln Met Pro Thr Gly Ile Glu Lys Glu Glu Ser
                125
                                    130
                                                         135
Glu His His Leu Pro Ala Gly Leu Phe Gln His Ser Arg Cys Met
                140
                                    145
                                                         150
Ala Arg Arg Gly Ile Tyr Met Val Ile Pro Val Pro Glu Ala Glu
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                                    160
Ser Asn Tyr Cys Leu Leu
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Pro Gly Met Ser Val Ala Gly Val Glu Gly Glu Pro Leu Val Ser
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Ser Gln Ser Gly Gln Ser Pro Pro Glu Pro Gln Asp Pro Glu Ala
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Pro Ser Ser Gly Pro Gly His Leu Val Ala Met Gly Lys Val
                                      40
Ser Arg Thr Pro Val Glu Ala Gly Val Ser Gln Ser Asp Ala Glu
                                      55
Asn Ala Ala Pro Ser Cys Pro Asp Glu His Asp Thr Leu Pro Arg
                 65
                                      70
                                                          75
Arg Arg Gly Arg Pro Ser Arg Arg Phe Leu Gly Lys Lys Tyr Arg
                 80
                                      85
Lys Tyr Tyr Tyr Lys Ser Pro Lys Pro Leu Leu Arg Pro Phe Leu
                 95
                                    100
Cys Arg Ile Cys Gly Ser Arg Phe Leu Ser His Glu Asp Leu Arg
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Phe His Val Asn Ser His Glu Ala Gly Asp Pro Gln Leu Phe Lys
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                                     130
                                                         135
Cys Leu Gln Cys Ser Tyr Arg Ser Arg Arg Trp Ser Ser Leu Lys
                140
Glu His Met Phe Asn His Val Gly Ser Lys Pro Tyr Lys Cys Asp
                155
                                     160
                                                          165
Glu Cys Ser Tyr Thr Ser Val Tyr Arg Lys Asp Val Ile Arg His
                170
                                     175
                                                          180
Ala Ala Val His Ser Arg Asp Arg Lys Lys Arg Pro Asp Pro
                                     190
                                                          195
                185
Pro Lys Leu Ser Ser Phe Pro Cys Pro Val Cys Gly Arg Val
                                                         Tvr
                200
                                     205
                                                          210
Pro Met Gln Lys Arg Leu Thr Gln His Met Lys Thr His Ser Thr
                215
                                     220
                                                          225
Glu Lys Pro His Met Cys Asp Lys Cys Gly Lys Ser Phe Lys Lys
                                     235
                230
Arg Tyr Thr Phe Lys Met His Leu Leu Thr His Ile Gln Ala Val
                                                          255
                245
                                     250
Ala Asn Arg Arg Phe Lys Cys Glu Phe Cys Glu Phe Val Cys Glu
                260
                                     `265
                                                          270
Asp Lys Lys Ala Leu Leu Asn His Gln Leu Ser His Val Ser Asp
                275
                                     280
                                                          285
Lys Pro Phe Lys Cys Ser Phe Cys Pro Tyr Arg Thr Phe Arg Glu
                                                          300
                290
                                     295
Asp Phe Leu Leu Ser His Val Ala Val Lys His Thr Gly Ala Lys
                305
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                                                          315
Pro Phe Ala Cys Glu Tyr Cys His Phe Ser Thr Arg His Lys Lys
                                     325
                320
Asn Leu Arg Leu His Val Arg Cys Arg His Ala Ser Ser Phe Glu
                335
                                     340
                                                          345
Glu Trp Gly Arg Arg His Pro Glu Glu Pro Pro Ser Arg Arg Arg
                                     355
                350
                                                          360
Pro Phe Phe Ser Leu Gln Gln Ile Glu Glu Leu Lys Gln Gln His
                                     370
                                                          375
                365
Ser Ala Ala Pro Gly Pro Pro Pro Ser Ser Pro Gly Pro Pro Glu
                380
                                     385
                                                          390
Ile Pro Pro Glu Ala Thr Thr Phe Gln Ser Ser Glu Ala Pro Ser
                395
Leu Leu Cys Ser Asp Thr Leu Gly Gly Ala Thr Ile Ile Tyr Gln
                410
                                     415
                                                          420
Gln Gly Ala Glu Glu Ser Thr Ala Met Ala Thr Gln Thr Ala Leu
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Asp Leu Leu Asn Met Ser Ala Gln Arg Glu Leu Gly Gly
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Phe Glu Asp Val Ala Val Asn Phe Thr Gln Glu Glu Trp Ala Leu
                                                           30
Leu Asp Ser Ser Gln Lys Asn Leu Tyr Arg Glu Val Met Gln Glu
                 35
                                      40
                                                           45
Thr Cys Arg Asn Leu Ala Ser Val Gly Ser Gln Trp Lys Asp Gln
                 50
                                      55
                                                           60
Asn Ile Glu Asp His Phe Glu Lys Pro Gly Lys Asp Ile Arg Asn
                                      70
                 65
His Ile Val Gln Arg Leu Cys Glu Ser Lys Glu Asp Gly Gln Tyr
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85
                 80
Gly Glu Val Val Ser Gln Ile Pro Asn Leu Asp Leu Asn Glu Asn
                 95
                                    100
                                                         105
Ile Ser Thr Gly Leu Lys Pro Cys Glu Cys Ser Ile Cys Gly Lys
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                                     115
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Val Phe Val Arg His Ser Leu
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Asn Ser Leu Ser Val Ala Ser Ala Pro Pro Gln Arg Asp Pro Gly
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Met Ala Met Ala Leu Pro Met Pro Gly Pro Gln Glu Ala Val Val
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                                                          30
Phe Glu Asp Val Ala Val Tyr Phe Thr Arg Ile Glu Trp Ser Cys
                                                           45
                 35
                                      40
Leu Ala Pro Asp Gln Gln Ala Leu Tyr Arg Asp Val Met Leu Glu
                 50
                                      55
                                                          60
Asn Tyr Gly Asn Leu Ala Ser Leu Gly Phe Leu Val Ala Lys Pro
                                      70
                 65
Ala Leu Ile Ser Leu Leu Glu Gln Gly Glu Glu Pro Gly Ala Leu
                                                           90
                 80
                                      85
Ile Leu Gln Val Ala Glu Gln Ser Val Ala Lys Ala Ser Leu Cys
                                     100
                                                          105
                 95
Thr Glu Asp Pro Asn Thr Leu Pro Ser Arg Ser Gln Glu Gly Ser
                110
                                     115
                                                          120
Pro Ala Ser Ser Glu Gly Gly Pro Gly Glu Lys Gly Val Ala Gly
                                                          135
                125
                                     130
Xaa Val Ala Gly Gly Gly Ala Ala Ser Ser Trp Pro His Gly Glu
                                     145
                                                          150
                140
His Pro Val Thr Pro Asn Arg
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Asp Thr Met Gln Ala Val Val Pro Leu Asn Lys Met Thr Ala Ile
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Ser Pro Glu Pro Gln Thr Leu Ala Ser Thr Glu Gln Asn Glu Val
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                                      25
Pro Arg Val Val Thr Ser Gly Glu Glu Ala Ile Leu Arg Gly
                                                           45
                                      40
                 35
Asn Ala Ala Asp Ala Glu Ser Phe Arg Gln Arg Phe Arg Trp Phe
                                      55
                 50
                                                           60
Cys Tyr Ser Glu Val Ala Gly Pro Arg Lys Ala Leu Ser Gln Leu
Trp Glu Leu Cys Asn Gln Trp Leu Arg Pro Asp Ile His Thr Lys
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85
Glu Gln Ile Leu Glu Leu Leu Val Phe Glu Gln Phe Leu Thr Ile
                 95
                                     100
                                                          105
Leu Pro Gly Glu Ile Arg Ile Trp Val Lys Ser Gln His Pro Glu
                                     115
                                                          120
                 110
Ser Ser Glu Glu Val Val Thr Leu Ile Glu Asp Leu Thr Gln Met
                                     130
                125
                                                          135
Leu Glu Glu Lys Asp Pro Val Ser Gln Asp Ser Thr Val Ser Gln
                                     145
                                                          150
                140
Glu Glu Asn Ser Lys Glu Asp Lys Met Val Thr Val Cys Pro Asn
                                                          165
                                     160
                155
Thr Glu Ser Cys Glu Ser Ile Thr Leu Lys Asp Val Ala Val Asn
                170
                                     175
Phe Ser Arg Gly Glu Trp Lys Lys Leu Glu Pro Phe Gln Lys Glu
                                     190
                185
                                                          195
Leu Tyr Lys Glu Val Leu Leu Glu Asn Leu Arg Asn Leu Glu Phe
                                                          210
                200
                                     205
Leu Asp Phe Pro Val Ser Lys Leu Glu Leu Ile Ser Gln Leu Lys
                                     220
                                                          225
                215
Trp Val Glu Leu Pro Trp Leu Leu Glu Glu Val Ser Lys Ser Ser
                230
                                     235
                                                          240
Arg Leu Asp Glu Ser Ala Leu Asp Lys Ile Ile Glu Arg Cys Leu
                                                          255
                245
                                     250
Arg Asp Asp His Gly Leu Met Glu Glu Ser Gln Gln Tyr Cys
                                     265
                260
Gly Ser Ser Glu Glu Asp His Gly Asn Gln Gly Asn Ser Lys Gly
                                                          285
                275
                                     280
Arg Val Ala Gln Tyr Lys Thr Leu Gly Ser Gly Ser Arg Gly
                                                         Lvs
                 290
                                     295
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Lys Phe Asp Pro Asp
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Pro Lys Gly Gln Gly Ser Gln Glu His Leu Leu Lys Leu Met Thr
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                                      25
                                                           30
                     Thr Pro Asp Phe Asp Asp Leu Leu Ala Ala
Met Gly Asp Met Lys
                  35
                                      40
                                                           45
Phe Asp Ile Pro Asp Met Val Asp Pro Lys Ala Ala Ile Glu Ser
                                      55
                                                           60
                  50
Gly His Asp Asp His Glu Ser His Met Lys Gln Asn Ala His Gly
                                      70
                  65
Glu Asp Asp Ser His Ala Pro Ser Ser Asp Val Gly Val Ser
                                      85
                                                           90
                  80
                                         Ser Ser Glu Gly Gly
Val Ile Val Lys Asn Val Arg Asn Ile Asp
                  95
                                     100
                                                          105
Glu Lys Asp Gly His Asn Pro Thr Gly Asn Gly Leu His Asn Gly
                                                          120
                 110
                                     115
Phe Leu Thr Ala Ser Ser Leu Asp Ser Tyr Ser Lys Asp Gly Ala
                                                          135
                                     130
                 125
Lys Ser Leu Lys Gly Asp Val Pro Ala Ser Glu Val Thr Leu Lys
                                                          150
                                     145
                 140
Asp Ser Thr Phe Ser Gln Phe Ser Pro Ile Ser Ser Ala Glu Glu
                 155
                                     160
                                                          165
Phe Asp Asp Asp Glu'Lys Ile Glu Val Asp Asp Pro Pro Asp Lys
                 170
                                     175
Glu Asp Met Arg Ser Ser Phe Arg Ser Asn Val Leu Thr Gly Ser
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185
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                                                          195
Ala Pro Gln Gln Asp Tyr Asp Lys Leu Lys Ala Leu Gly Gly Glu
                200
                                     205
                                                          210
Asn Ser Ser Lys Thr Gly Leu Ser Thr Ser Gly Asn Val Glu
                                                          Ьуs
                                     220
                                                          225
                215
Asn Lys Ala Val Lys Arg Glu Thr Glu Ala Ser Ser Ile Asn Leu
                                                          240
                230
                                     235
                                                          Leu
Ser Val Tyr Glu Pro Phe Lys Val Arg Lys Ala Glu Asp Lys
                245
                                     250
                                                          255
Lys Glu Ser Ser Asp Lys Val Leu Glu Asn Arg Val Leu Asp
                                                          Gly
                                                          270
                260
                                     265
Lys Leu Ser Ser Glu Lys Asn Asp Thr Ser Leu Pro Ser Val Ala
                275
                                     280
                                                          285
Pro Ser Lys Thr Lys Ser Ser Ser Lys Leu Ser Ser Cys Ile Ala
                290
                                     295
                                                          300
Ala Ile Ala Ala Leu Ser Ala Lys Lys Ala Ala Ser Asp Ser
                                                          Cys
                305
                                     310
                                                          315
Lys Glu Pro Val Ala Asn Ser Arg Glu Ser
                                         Ser Pro Leu Pro
                                                          Lys
                                                          330
                320
                                     325
Glu Val Asn Asp Ser Pro Arg Ala Ala Asp Lys Ser Pro Glu Ser
                                                          345
                335
                                     340
Gln Asn Leu Ile Asp Gly Thr Lys Lys Pro
                                         Ser Leu Lys Gln Pro
                350
                                     355
                                                          360
Asp Ser Pro Arg Ser Ile Ser Ser Glu Asn Ser Ser Lys Gly Ser
                                     370
                                                          375
                 365
Pro Ser Ser Pro Ala Gly Ser Thr Pro Ala Ile Pro Lys Val Arg
                 380
                                     385
                                                          390
Ile Lys Thr Ile Lys Thr Ser Ser Gly Glu
                                         Ile Lys Arg Thr Val
                                     400
                                                          405
                 395
Thr Arg Val Leu Pro Glu Val Asp Leu Asp
                                         Ser Gly Lys Lys Pro
                                                          420
                                     415
                 410
Ser Glu Gln Thr Ala Ser Val Met Ala Ser Val Thr Ser Leu Leu
                                     430
                 425
                                         Ser Pro Pro Arg Ala
Ser Ser Pro Ala Ser Ala Ala Val Leu Ser
                                      445
                                                           450
                 440
Pro Leu Gln Ser Ala Val Val Thr Asn Ala Val Ser Pro Ala Glu
                 455
                                      460
                                                           465
Leu Thr Pro Lys Gln Val Thr Ile Lys Pro Val Ala Thr Ala Phe
                                                           480
                 470
                                      475
Leu Pro Val Ser Ala Val Lys Thr Ala Gly Ser Gln Val Ile Asn
                                      490
                 485
Leu Lys Leu Ala Asn Asn Thr Thr Val Lys Ala Thr Val Ile Ser
                 500
                                      505
Ala Ala Ser Val Gln Ser Ala Ser Ser Ala Ile Ile Lys Ala Ala
                 515
                                      520
                                                           525
Asn Ala Ile Gln Gln Gln Thr Val Val Val Pro Ala Ser Ser Leu
                 530
                                      535
                                                           540
Ala Asn Ala Lys Leu Val Pro Lys Thr Val His Leu Ala Asn Leu
                                                           555
                 545
                                      550
Asn Leu Leu Pro Gln Gly Ala Gln Ala Thr Ser Glu Leu Arg Gln
                                                           570
                                      565
                 560
Val Leu Thr Lys Pro Gln Gln Gln Ile Lys Gln Ala Ile Ile Asn
                                      580
                                                           585
                 575
                                          Ser Arg Val Gln Val
Ala Ala Ala Ser Gln Pro Pro Lys Lys Val
                 590
                                      595
                                                           600
Val Ser Ser Leu Gln Ser Ser Val Val Glu Ala Phe Asn Lys Val
                 605
                                      610
                                                           615
Leu Ser Ser Val Asn Pro Val Pro Val Tyr Ile Pro Asn Leu Ser
                 620
                                      625
                                                           630
Pro Pro Ala Asn Ala Gly Ile Thr Leu Pro Thr Arg Gly Tyr Lys
                                                           645
                 635
                                      640
Cys Leu Glu Cys Gly Asp Ser Phe Ala Leu Glu Lys Ser Leu Thr
                                                           660
                                      655
                 650
Gln His Tyr Asp Arg Arg Ser Val Arg Ile Glu Val Thr
                                      670
                                                           675
                 665
His Cys Thr Lys Asn Leu Val Phe Tyr Asn Lys Cys Ser Leu Leu
                                      685
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Ser His Ala Arg Gly His Lys Glu Lys Gly Val Val Met Gln Cys

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695
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Ser His Leu Ile Leu Ser Gln Ser Gln Gln Ile Lys
                710
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Thr Pro Lys Pro Gln Val Ile Ser Leu Leu Glu Gln Gly Lys Glu
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Pro Trp Met Val Gly Arg Glu Leu Thr Xaa Gly Leu Cys Ser Asp
                                      25
                 20
Leu Glu Ser Met Cys Glu Thr Lys Leu Leu Ser Leu Lys Lys Glu
                                      40
Val Tyr Glu Ile Glu Leu Cys Gln Xaa Glu Ile Met Gly Leu Thr
                 50
                                      55
                                                           60
Lys His Gly Leu Glu Tyr Ser Ser Phe Gly Asp Val Leu Glu Tyr
                 65
                                      70
Arg Ser His Leu Ala Lys Gln Leu Gly Tyr Pro Asn Gly His Phe
                                      85
                                                           90
                 80
Ser Gln Xaa Ile Phe Thr Pro Glu Tyr Met Pro Thr Phe Ile Gln
                                     100
                                                          105
                 95
Gln Thr Phe Leu Thr Leu His Xaa Ile Ile Xaa Asn Glu Asp Arg
                                                          120
                110
                                     115
Xaa Tyr Glu Cys Lys Glu Cys Gly Lys Met Phe Ser His Gly Ser
                125
                                     130
                                                          135
Gln Leu Thr Gln His Gln Arg Ile His Thr Gly Glu Lys Pro Tyr
                140
                                     145
                                                          150
Gln Cys Lys Glu Cys Gly Lys Ala Phe Asn Arg Gly Ser Leu Leu
                                     160
                                                          165
                155
Thr Arg His Gln Arg Ile His Thr Gly Glu Lys Pro Tyr Glu Cys
                170
                                     175
                                                          180
Lys Glu Cys Gly Lys Thr Phe Ser Arg Gly Ser Glu Leu Thr Gln
                                     190
                                                          195
                185
His Glu Arg Ile His Thr Ala Gly Ala Pro Leu Leu Ser Trp Gly
                                     205
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Leu
<210> 255
<211> 103
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Gly Glu Gly Gln Ser Ala Asp Leu Arg Pro Leu Leu Thr Asp
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Phe Arg Leu Gln Phe Ile Cys Ala Pro Ala Ser Ser Leu Ser Leu
                                      25
                                                           30
                 20
Arg Arg Leu Arg Leu Arg Pro Gln Lys Glu Ile Ser Ile Leu Cys
                                                           45
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Pro Glu Gln Asn Arg Met Ala Met Ser Gln Glu Ser Leu Thr Phe
                 50
                                      55
Lys Asp Val Phe Val Gly Phe Thr Leu Glu Glu Trp Gln Gln Leu
                 65
                                      70
Asp Pro Ser Gln Arg Ala Leu Tyr Arg Asp Val Met Leu Glu Asn
                                      85
Tyr Ser Asn Leu Val Ser Val Gly Tyr Cys Ala His Lys
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                                     100
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<211> 84
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Glu Gly Ile Pro Glu Lys Lys Glu Glu Glu Glu Met Ala Gly
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Ser Gln Gly Leu Leu Ile'Phe Arg Asp Val Ala Ile Glu Phe Ser
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                                      25
Pro Glu Glu Trp Ser Tyr Leu Asp Pro Ala Gln Gln Asn Leu Tyr
                                      40
                                                           45
                 35
Arg Asp Val Met Leu Glu Asn Tyr Arg Asn Leu Val Ser Leu Gly
                 50
                                      55
                                                           60
Ile Ala Val Ser Lys Pro Glu Leu Ile Thr Cys Leu Glu Gln Arg
                 65
                                      70
Asn Glu Pro Trp Asn Val Lys Lys His
                 80
<210> 257
<211> 194
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Asp Ala Arg Thr Thr Trp Lys Pro Arg Asn Val Ile Tyr Ser His
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                                                           15
Phe Thr Glu Asp Leu Trp Pro Glu His Ser Ile Lys Asp Ser Phe
                 20
                                      25
Gln Lys Val Ile Leu Arg Gly Tyr Gly Lys Cys Gly His Glu Asn
                                      40
                                                           45
                 35
Leu Gln Leu Arg Ile Ser Cys Lys Ser Val Asp Glu Ser Lys Val
                                                           60
                 50
                                      55
Phe Lys Glu Gly Tyr Asn Glu Leu Asn Gln Cys Leu Arg Thr Thr
                 65
                                      70
Gln Ser Lys Ile Phe Gln Cys Asp Lys Tyr Val Lys Val Phe His
                 80
                                      85
                                                           90
Lys Phe Ser Asn Ser Asn Ser His Lys Lys Arg Asn Thr Gly Lys
                 95
                                     100
                                                          105
Lys Val Phe Lys Cys Lys Glu Cys Gly Lys Ser Phe Cys Met Leu
                                                          120
                110
                                     115
Ser His Leu Thr Gln His Ile Arg Ile His Thr Arg Glu Asn Ser
                                                          135
                125
                                     130
Tyr Lys Cys Lys Glu Cys Gly Lys Val Leu Asn Gln Ser Ser Glu
                140
                                     145
                                                          150
Leu Ile Lys His Lys Lys Ile His Thr Gly Glu Lys Pro Tyr Thr
                155
                                     160
Cys Glu Lys Cys Gly Lys Thr Phe Asn Gln Ser Ala Asn Leu Tyr
                170
                                     175
Ala His Lys Lys Ile His Thr Gly Asp Lys Thr Ile Gln Val
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PCT/US01/06059

WO 01/62927 185 190 <210> 258 <211> 129 <212> PRT <213> Homo sapiens <220> <221> misc\_feature <223> Incyte ID No: LG:1082586.1.orf1:2000FEB18 <221> unsure <222> 5 <223> unknown or other <400> 258 Ala Gln Pro Arg Xaa Pro Met Gly Gln Tyr Gln Ala Asp Gln Tyr 10 Ile His Arg Arg Ser Ser Ser Arg Arg Glu Lys Gly Ala Glu Arg 20 25 3.0 Ile Leu Glu Glu Ile Met Ala Glu Asn Phe Ser Ser Leu Ile Lys 35 40 Asp Met Asn Ile Asn Ile Gln Glu Ala Gln Gln Thr Pro Ser Met 55 60 50 Met Asn Ser Lys Ile Ala Thr Leu Arg His Ile Ile Lys Leu 70 65 Ser Lys Asp Lys His Arg Pro Leu Ser Leu Thr Ala Ala Arg Ala 85 90 80 Pro Ser Leu Val Phe Thr Ser Leu Phe Leu Leu Leu Glu Ala 95 100 105 Gln Pro Leu Trp Pro Cys Asp Leu Gln Val Leu Gly Asp Pro Leu 110 115 120 Leu Arg Cys Gln Asp Pro Leu Glu Ala 125 <210> 259 <211> 93 <212> PRT <213> Homo sapiens <220> <221> misc\_feature <223> Incyte ID No: LG:1082774.1.orf3:2000FEB18 <400> 259 Pro Ala Gly Ile Arg Arg Val Thr Ala Arg Thr Pro Gly Pro Pro 1 10 15 Gly Ser Leu Glu Met Gly Pro Leu Gln Phe Arg Asp Val Ala Ile 25 30 20 Glu Phe Ser Leu Glu Glu Trp His Cys Leu Asp Ala Ala Gln Arg 45 40 35 Asn Leu Tyr Arg Asp Val Met Leu Glu Asn Tyr Arg Asn Leu Ile 55 50 Phe Leu Gly Ile Val Val Ser Lys Pro Asn Leu Ile Thr Cys Leu 70 65 Glu Gln Gly Lys Lys Pro Leu Thr Met Lys Arg His Glu Met Ile

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Ala Lys Pro

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80

85

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Lys His Glu Ile Ile His Phe Glu Glu Glu Pro Ser Glu Tyr Asn
Asn Asn Gly Asn Ser Phe Trp Leu Asn Glu Asp Leu Ile Trp His
                 20
                                      25
                                                           30
Gln Lys Ile Lys Asn Trp Glu Gln Pro Phe Glu Tyr Asn Glu Cys
                 35
                                      40
                                                           45
Gly Lys Ala Phe Pro Glu Asn Ser Leu Phe Leu Val His Lys Arg
                 50
                                      55
                                                           60
Ala Tyr Thr Gly Gln Lys Thr Cys Lys Tyr
                                         Thr Glu His Gly Lys
                 65
Thr Cys Tyr Met Ser Phe Phe Ile Thr His Gln Gln Thr His Pro
                 80
                                      85
                                                           90
Arg Glu Asn His Tyr Glu Cys Asn Glu Cys Gly Glu Ser Ile Phe
                 95
                                     100
Glu Glu Ser Ile Leu Phe Glu His Gln Asn Val Tyr Pro Phe Ser
                                                          120
                110
                                     115
Gln Asn Leu Asn Pro Thr Leu Ile Gln Arg Thr His Ser Ile Ser
                125
                                     130
                                                          135
Asn Ile Ile Glu Tyr Asn Glu Cys Gly Thr Phe Phe Ser Glu Lys
                140
                                     145
                                                          150
Leu Ala Leu His Leu Gln Gln Arg Thr His Pro Gly Glu Lys
                155
                                     160
                                                          165
Tyr Glu Cys His Glu Cys Gly Lys Thr Phe Thr Gln Lys Ser Ala
                170
                                     175
His Thr Arg His Gln Arg Thr His Thr Gly Lys Thr Leu
                185
                                     190
<210> 261
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Pro Gly Leu Arg Asp Leu Thr Cys Lys Glu Leu Leu Ile Leu Thr
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Glu Arg Glu Ala Gln Lys Arg Lys Lys Arg Lys Glu Lys Glu Ser
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                                                           30
Gly Met Ala Leu Thr Gln Gly Pro Leu Thr Phe Arg Asp Val Ala
                                      40
Ile Glu Phe Ser Gln Glu Glu Trp Lys Ser Leu Asp Pro Val Gln
                                      55
                 50
                                                           60
Lys Ala Leu Tyr Trp Asp Val Met Leu Glu Asn Tyr Arg Asn Leu
                                      70
                                                           75
                 65
Val Phe Leu Gly Lys Asp Asn Phe Ala Leu Glu Val Lys Ile
                                                          Cys
                 80
                                      85
Pro Arg Val Phe Leu Tyr Phe Leu Cys Cys Leu Ser Val Gly Ala
                 95
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Arg Ser Ile His Leu His
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<210> 262
<211> 137
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Leu His His Ser Pro Cys Tyr Pro Val Thr Cys Arg Tyr Trp Asp
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Ile His Arg Glu Glu Gly Gly Thr Ser Gly Gly Trp Glu Met
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                                                           30
Val Leu Thr Phe Arg Asp Val Ala Val Glu Phe Ser Pro Glu Glu
                                      40
Trp Glu Cys Leu Asp Ser Ala Gln Gln Arg Leu Tyr Arg Asp Val
                 50
                                      55
Met Leu Glu Asn Tyr Gly Asn Leu Phe Ser Leu Gly Leu Ala Ile
                                      70
                                                           75
Phe Lys Pro Asp Leu Ile Thr Tyr Leu Glu Gln Arg Lys Glu Pro
                 80
                                      85
                                                           90
Trp Asn Ala Arg Arg Gln Lys Thr Val Ala Lys His Pro Asp Tyr
                 95
                                     100
                                                          105
Tyr Asp Val Cys Asn Glu Asp Tyr Glu Tyr Asn Trp Ser Tyr Met
                                     115
                                                          120
                110
Phe Leu Asn Ser Glu Gln Leu Phe Ile Lys Phe Tyr Pro Thr Phe
                                     130
                125
Phe Cys
<210> 263
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Met Phe Pro Val Leu Glu Pro His Gln Val Gly Leu Ile Arg Ser
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                  5
Tyr Asn Ser Lys Thr Met Thr Cys Phe Gln Glu Leu Val Thr Phe
                                                           30
                  20
Arg Asp Val Ala Ile Asp Phe Ser Arg Gln Glu Trp Glu Cys Leu
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Asp Pro Asn Gln Arg Asp Leu Tyr Arg Asp Val Met Leu Glu Asn
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Tyr Arg Asn Leu Val Ser Leu Gly
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<210> 264
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Asp Thr Gly Thr Ser Trp Lys Pro Lys Met Gly Pro Leu Gln Phe
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Arg Asp Val Ala Ile Glu Phe Ser Leu Glu Glu Trp His Cys Leu
                                                           30
                  20
                                      25
Asp Thr Ala Gln Arg Asn Leu Tyr Arg Asn Val Met Leu Glu Asn
                                      40
                                                           45
Tyr Ser Asn Leu Val Phe Leu Gly Ile Thr Val Ser Lys Pro Asp
                                      55
                                                           60
                  50
                                                          Lys
75
Leu Ile Thr Cys Leu Glu Gln Gly Arg Lys Pro Leu Thr Met
                                       70
                  65
Arg Asn Glu Met Ile Ala Lys Pro Ser Val Ser Phe Leu Gln Val
                  80
                                      85
                                                           90
His Ser Glu Ser Gln Ser Pro Leu His Asp Ile
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                  95
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<211> 96
<212> PRT
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Ser Ala Ala Met Phe Pro Val Phe Ser Gly Cys Phe Gln Glu Leu
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Gln Glu Lys Asn Lys Ser Leu Glu Leu Val Ser Phe Glu Glu Val
                 20
                                      25
                                                           3.0
Ala Val His Phe Thr Trp Glu Glu Trp Gln Asp Leu Asp Asp Ala
                 35
                                      40
                                                           45
Gln Arg Thr Leu Tyr Arg Asp Val Met Leu Glu Thr Tyr Ser Ser
                                      55
                                                           60
                 50
Leu Val Ser Leu Gly His Cys Ile Thr Lys Pro Glu Met Ile Phe
                                      70
                 65
Lys Leu Glu Gln Gly Ala Glu Pro Trp Ile Val Glu Glu Thr Leu
                 8.0
Asn Leu Arg Leu Ser Gly
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<400> 266
Glu Asn Gly Glu Asn Cys Asn Gln Asp Met Phe Glu Asn Glu Ser
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Arg Lys Ile Phe Ser Glu Met Pro Glu Gly Glu Ser Ala Gln His
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Ser Asp Gly Glu Ser Asp Phe Glu Arg Asp Ala Gly Ile Gln Arg
                                      40
                                                           45
Leu Gln Gly His Thr Pro Gly Glu Asp His Gly Glu Val Val Ser
                 50
                                      55
                                                           60
Gln Asp Arg Glu Val Gly Gln Leu Ile Gly Leu Gln Gly Thr Tyr
                                      70
                 65
Leu Gly Glu Lys Pro Tyr Glu Cys Pro Gln Cys Gly Lys Thr Phe
                                      85
                                                           90
                 80
Ser Arg Lys Ser His Leu Ile Thr His Glu Arg Thr His Thr Gly
                                                          105
                 95
                                     100
Glu Lys Tyr Tyr Lys Cys Asp Glu Cys Gly Lys Ser Phe Ser Asp
                                     115
                                                          120
                 110
Gly Ser Asn Phe Ser Arg His Gln Thr Thr His Thr Gly Glu Lys
                125
                                     130
                                                          135
Pro Tyr Lys Cys Arg Asp Cys Gly Lys Ser Phe Ser Arg Ser Ala
                 140
                                     145
                                                          150
Asn Leu Ile Thr His Gln Arg Ile His Thr Gly Glu Lys Pro Phe
                                                          165
                 155
                                     160
Gln Cys Ala Glu Cys Gly Lys Ser Phe Ser Arg Ser Pro Asn Leu
                 170
                                     175
                                                          180
Ile Ala His Gln Arg Thr His Thr Gly Glu Lys Pro Tyr Ser Cys
                 185
                                     190
Pro Glu Cys Gly Lys Ser Phe Gly Asn Arg Ser Ser Leu Asn Thr
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200
                                     205
                                                          210
His Gln Gly Ile His Thr Gly Glu Lys Pro Tyr Glu Cys Lys Glu
                215
                                     220
                                                          225
Cys Gly Glu Ser Phe Ser Tyr Asn Xaa Asn Leu Ile Arg Gln Gln
                230
                                     235
Arg Ile His Thr Gly Glu Lys Pro Tyr Lys
                245
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Pro Gly Met Ser Val Ala Gly Val Glu Gly Glu Pro Leu Val Ser
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Ser Gln Ser Gly Gln Ser Pro Pro Glu Pro Gln Asp Pro Glu Ala
                 20
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Pro Ser Ser Gly Pro Gly His Leu Val Ala Met Gly Lys Val
                                                           45
                                      40
Ser Arg Thr Pro Val Glu Ala Gly Val Ser Gln Ser Asp Ala Glu
                 50
                                      55
                                                           60
Asn Ala Ala Pro Ser Cys Pro Asp Glu His Asp Thr Leu Pro Arg
                 65
                                      70
                                                           75
Arg Arg Gly Arg Pro Ser Arg Arg Phe Leu Gly Lys Lys
                                                     Tyr Arg
                                                           90
                                      85
                 80
                    Ser Pro Lys Pro Leu Leu Arg Pro Phe Leu
Lys Tyr Tyr Tyr Lys
                 95
                                                          105
                                     100
Cys Arg Ile Cys Gly Ser Arg Phe Leu Ser His Glu Asp Leu Arg
                                     115
                                                          120
                110
Phe His Val Asn Ser His Glu Ala Gly Asp Pro Gln Leu Phe Lys
                 125
                                     130
                                                          135
Cys Leu Gln Cys Ser Tyr Arg Ser Arg Arg Trp Ser Ser Leu Lys
                140
                                     145
                                                          150
Glu His Met Phe Asn His Val Gly Ser Lys Pro Tyr Lys Cys Asp
                 155
                                     160
                                                          165
Glu Cys Ser Tyr Thr Ser Val Tyr Arg Lys Asp Val Ile Arg His
                 170
                                     175
                                                          180
Ala Ala Val His Ser Arg Asp Arg Lys Lys Arg Pro Asp Pro Thr
                                                          195
                 185
                                     190
Pro Lys Leu Ser Ser Phe Pro Cys Pro Val Cys Gly Arg Val Tyr
                 200
                                     205
                                                          210
Pro Met Gln Lys Arg Leu Thr Gln His Met Lys Thr His Ser Thr
                 215
                                     220
                                                          225
Glu Lys Pro His Met Cys Asp Lys Cys Gly Lys Ser Phe Lys Lys
                                     235
                                                          240
                 230
Arg Tyr Thr Phe Lys Met His Leu Leu Thr His Ile Gln Ala Val
                                     250
                                                          255
                 245
Ala Asn Arg Arg Phe Lys Cys Glu Phe Cys Glu Phe Val Cýs Glu
                                     265
                 260
                                                          270
Asp Lys Lys Ala Leu Leu Asn His Gln Leu Ser His Val Ser Asp
                 275
                                      280
                                                           285
Lys Pro Phe Lys Cys Ser Phe Cys Pro Tyr Arg Thr Phe Arg Glu
                 290
                                     295
                                                          300
Asp Phe Leu Leu Ser His Val Ala Val Lys His Thr Gly Ala Lys
                 305
                                     310
                                                          315
Pro Phe Ala Cys Glu Tyr Cys His Phe Ser Thr Arg His Lys Lys
                                                          330
                                     325
                 320
Asn Leu Arg Leu His Val Arg Cys Arg His Ala Ser Ser Phe Glu
                 335
                                     340
                                                          345
Glu Trp Gly Arg Arg His Pro Glu Glu Pro Pro Ser Arg Arg Arg
                 350
                                      355
Pro Phe Phe Ser Leu Gln Gln Ile Glu Glu Leu Lys Gln Gln His
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370
                365
                                                         375
Ser Ala Ala Pro Gly Pro Pro Pro Ser Ser Pro Gly Pro Pro Glu
                380
                                     385
                                                          390
Ile Pro Pro Glu Ala Thr Thr Phe Gln Ser Ser Glu Ala Pro Ser
                395
                                     400
Leu Leu Cys Ser Asp Thr Leu Gly Gly Ala Thr Ile Ile Tyr Gln
                410
                                     415
                                                          420
Gln Gly Ala Glu Glu Ser Thr Ala Met Ala Thr Gln Thr Ala Leu
                425
                                     430
                                                          435
Asp Leu Leu Asn Met Ser Ala Gln Arg Gly Pro Gly Gly Thr
                440
                                     445
                                                          450
Ala Leu Gln Val Cys Cys Leu Gly Thr Cys Ser Pro Ser Gln Leu
                455
                                     460
Pro Gln Tyr Pro Ala Leu His Trp Thr Leu Gly Leu Glu Glu Asn
                                     475
                                                          480
                470
Ser Val Ser Glu Leu Leu Arg Pro Trp Gly Leu Pro Gly Ser Gly
                                     490
                                                          495
                485
Gly Asp Arg Ser Ala Glu Val Trp Trp Ala Asn Arg Glu Glu Gln
                500
                                     505
Ala Leu Pro Arg Arg Pro Gln Gly Ile Pro Ser Ile
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<222> 36
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Gly Leu Gly Glu Glu Val Pro Cys Ala Met Met Glu Gly Val Ala
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Ala Tyr Thr Gln Thr Glu Pro Glu Gly Ser Gln Pro Ser Thr Met
                                                           30
                 20
Asp Ala Thr Ala Val Xaa Gly Ile Glu Thr Lys Lys Glu Lys Glu
                 35
                                      40
                                                           45
Asp Leu Cys Leu Leu Lys Lys Glu Glu Lys Glu Glu Pro Val Ala
                                                           60
                 50
                                      55
Pro Glu Leu Ala Thr Thr Val Pro Glu Ser Ala Glu Pro Glu Ala
                                                           75
                                      70
                 65
Glu Ala Asp Gly Glu Glu Leu Asp Gly Ser Asp Met Ser Ala Ile
                                      85
                 80
                                                         Lvs
Ile Tyr Glu Ile Pro Lys Glu Pro Glu Lys Arg Arg Arg Ser
                                                          105
                 95
                                     100
Arg Ser Arg Val Met Asp Ala Asp Gly Leu Leu Glu Met Phe His
                110
                                     115
                                                          120
Cys Pro Tyr Glu Gly Cys Ser Gln Val Tyr Val Ala Leu Ser Ser
                125
                                     130
                                                          135
Phe Gln Asn His Val Asn Leu Val His Arg Lys Gly Lys Thr Lys
                                                          150
                140
                                     145
Val Cys Pro His Pro Gly Cys Gly Lys Lys Phe Tyr Leu Ser Asn
                                                          165
                155
                                     160
His Leu Arg Arg His Met Ile Ile His Ser Gly Val Arg Glu Phe
                                     175
                                                          180
                170
Thr Cys Glu Thr Cys Gly Lys Ser Phe Lys Arg Lys Asn His Leu
                                     190
                                                          195
                185
Glu Val His Arg Arg Thr His Thr Gly Glu Thr Pro Leu Gln Cys
                                     205
                 200
                                                          210
Glu Ile Cys Gly Tyr Gln Cys Arg Gln Arg Ala Ser Leu Asn Trp
                                                          225
                                     220
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His Met Lys Lys His Thr Ala Glu Val Gln Tyr Asn Phe Thr Cys
                230
                                     235
                                                          240
Asp Arg Cys Gly Lys Arg Phe Glu Lys Leu Asp Ser Val Lys Phe
                                     250
                245
                                                          255
His Thr Leu Lys Ser His Pro Asp His Lys Pro Thr
                260
                                     265
<210> 269
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<212> PRT
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Arg Pro Leu Pro Ala Asp Leu Pro Val Gly Gly His His Cys Leu
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His Gly Pro Gln Glu Ala Gly Leu Ser Ala Leu Gln Arg Pro Gln
                 20
                                      25
                                                           30
Pro Arg Pro Gly Leu Arg Thr Arg Gly Ala Glu Gly Leu Glu Leu
                                                           45
                                      40
                 35
Pro Ala Leu Trp Gln Thr Val His Ser Gly Leu Glu Ala Ala Ala
                                                           60
                 50
Ser Arg Pro Val Gly Pro Arg Thr Val His Leu Pro Asp Arg Ile
Arg Gly Pro Gly Gly Pro Val Leu Gly Leu Ala Glu Val Ala Ala
                 80
                                      85
                                                           9.0
Ala Val Ser Ala Val Val Gly Pro Ala Ala Glu Ala Lys Ser Pro
                                                          105
                 95
                                     100
Arg Ala Ser Gly Ser Gly Leu Thr Arg Arg Ser Pro Pro Val Leu
                                                          120
                110
                                     115
Cys Ala Arg Arg Pro Ser Ala Pro Ser Ala Thr Ser Lys Cys Thr
                                     130
                                                          135
                 125
Cys Ala His Thr Gln Ala Ser Gly Pro Met Leu Ala Thr Ser Val
                 140
                                     145
                                                          150
Pro Thr Pro Ala Pro Arg Ala Ala Ser Ser Thr Ala Thr Arg Arg
                 155
                                     160
                                                          165
Pro Thr Gly Arg Cys Arg Pro Arg Ala Pro Ser Trp Pro Thr Pro
                                                          180
                 170
                                     175
Ala Arg Ser Arg Pro Leu Gln Pro Leu Arg Ser Arg Leu Ser Met
                                                          195
                                     190
                 185
Leu Leu Pro Pro Pro Ala Pro Phe His Ala Ala Val Val Arg Gly
                                     205
                                                          210
                 200
Leu Glu Pro Pro Pro Gln Gln Val Ser Arg Asn Pro Gly Leu Leu
                                     220
                 215
Ala Val Gly Leu Lys Pro Ala Leu Val Glu Thr Leu Gly Glu Pro
                 230
                                     235
                                                          240
Ser Pro Arg Asn Lys Glu Leu Thr Leu Gln Thr Ala Arg Arg His
                                                          255
                 245
                                     250
His Pro Lys Arg Cys Pro Ser Gln Gly Ala Arg Ala Ala Gly Pro
                                                          270
                 260
                                     265
Gly Ala Ala Val Ser Ser Ala Gly Ser Ile Leu Pro Thr Ala Ala
                 275
                                     280
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<211> 194
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Gly Leu Ser Pro Lys Ala Ala Asn Leu Ala Pro Thr Thr Gln Gln
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Arg Ser Val Val Phe Pro Gln Thr Pro Cys Ser Arg Asn Phe Ser
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Leu Leu Asp Lys Ser Gly Pro Ile Glu Ser Gly Phe Asn Gln Ile
                 35
                                      40
Asn Val Lys Asn Gln Arg Val Leu Ala Ser Pro Thr Ser Thr Ser
                                      55
Gln Leu His Ser Glu Phe Ser Asp Trp His Leu Trp Lys Cys Gly
                                      70
                 65
Gln Cys Phe Lys Thr Phe Thr Gln Arg Ile Leu Leu Gln Met His
                                      85
                                                           90
                 80
Val Cys Thr Gln Asn Pro Asp Arg Pro Tyr Gln Cys Gly His Cys
                 95
                                     100
                                                          105
Ser Gln Ser Phe Ser Gln Pro Ser Glu Leu Arg Asn His Val Val
                110
                                     115
                                                          120
Thr His Ser Ser Asp Arg Pro Phe Lys Cys Gly Tyr Cys Gly Arg
                                     130
                                                          135
                125
Ala Phe Ala Gly Ala Thr Thr Leu Asn Asn His Ile Arg Thr His
                140
                                     145
Thr Gly Glu Lys Pro Phe Lys Cys Glu Arg Cys Glu Arg Ser Phe
                155
                                     160
                                                          165
Thr Gln Ala Thr Gln Leu Ser Arg His Gln Arg Met Pro Asn Glu
                170
                                     175
                                                          180
Cys Lys Pro Ile Thr Glu Ser Pro Glu Ser Ile Glu Val Asp
                185
                                     190
<210> 271
<211> 263
<212> PRT
<213> Homo sapiens
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<223> Incyte ID No: LI:758502.1.orf3:2000FEB01
Thr Leu Ile Lys His Gln Arg Thr His Thr Gly Glu Arg Pro Tyr
Glu Cys Pro Glu Cys Gly Lys Thr Phe Gly Arg Lys Pro His Leu
                 20
                                      25
                                                           30
Ile Met His Gln Arg Thr His Thr Gly Glu Lys Pro Tyr Ala Cys
                                                           45
                                      40
                 35
Leu Glu Cys His Lys Ser Phe Ser Arg Ser Ser Asn Phe Ile Thr
                                      55
                  50
                                                           60
His Gln Arg Thr His Thr Gly Val Lys Pro Tyr Arg Cys Asn Asp
                  65
                                      70
Cys Gly Glu Ser Phe Ser Gln Ser Ser Asp Leu Ile Lys His Gln
                 80
                                      85
                                                           90
Arg Thr His Thr Gly Glu Arg Pro Phe Lys Cys Pro Glu Cys Gly
                 95
                                     100
                                                          105
Lys Gly Phe Arg Asp Ser Ser His Phe Val Ala His Met Ser Thr
                                                          120
                110
                                     115
His Ser Gly Glu Arg Pro Phe Ser Cys Pro Asp Cys His Lys Ser
                                     130
                                                          135
                125
Phe Ser Gln Ser Ser His Leu Val Thr His Gln Arg Thr His Thr
                                                          150
                140
                                     145
Gly Glu Arg Pro Phe Lys Cys Glu Asn Cys Gly Lys Gly Phe Ala
                155
                                     160
Asp Ser Ser Ala Leu Thr Lys His Gln Arg Ile His Thr Gly Glu
                 170
                                     175
Arg Pro Tyr Lys Cys Gly Glu Cys Gly Lys Ser Phe Asn Gln Ser
                 185
                                     190
                                                          195
Ser His Phe Ile Thr His Gln Arg Ile His Leu Gly Asp Arg Pro
                200
                                     205
                                                          210
Tyr Arg Cys Pro Glu Cys Gly Lys Thr Phe Asn Gln Arg Ser His
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215
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                                                          225
Phe Leu Thr His Gln Arg Thr His Thr Gly Glu Lys Pro Phe His
                230
                                     235
                                                          240
Cys Ser Lys Cys Asn Lys Ser Phe Arg Gln Lys Ala His Leu Leu
                                     250
                245
Cys His Gln Asp Thr His Leu Ile
                260
<210> 272
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<222> 142
<223> unknown or other
<400> 272
Ala Lys Asn Leu Phe Lys Met Asp Ile Glu Asp Cys Asn Gly Arg
                                      10
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Ser Tyr Val Ser Gly Ser Gly Asp Ser Ser Leu Glu Lys Glu Phe
                                      25
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                  20
Leu Gly Ala Pro Val Gly Pro Ser Val Ser Thr Pro Asn Ser Gln
                 35
                                      40
His Ser Ser Pro Ser Arg Ser Leu Ser Ala Asn Ser Ile Lys Val
                  50
                                      - 55
                                                           60
Glu Met Tyr Ser Asp Glu Glu Ser Ser Arg Leu Leu Gly Pro Asp
                  65
                                      70
                                                           75
Glu Arg Leu Leu Glu Lys Asp Asp Ser Val Ile Val Glu Asp Ser
                                                           90
                  80
                                      85
Leu Ser Glu Pro Leu Gly Tyr Cys Asp Gly Ser Gly Pro Glu Pro
                  95
                                     100
                                                          105
His Ser Pro Gly Gly Ile Arg Leu Pro Asn Gly Lys Leu Lys
                                                          Cys
                                                          120
                 110
                                     115
Asp Val Cys Gly Met Val Cys Ile Gly Pro Asn Val Leu Met Val
                 125
                                     130
His Lys Arg Ser His Thr Xaa
                 140
<210> 273
<211> 164
<212> PRT
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<223> Incyte ID No: LI:789445.1.orf2:2000FEB01
<400> 273
Glu His Arg Glu Ala Lys Ala Ser Gly Trp Val Thr Asp Gly Leu
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Leu Met Asp Ser Ser Gln His Leu Val Thr Phe Glu Asp Val Ala
                                       25
                                                           3.0
Val Asp Phe Thr Gln Glu Glu Trp Thr Leu Leu Asp Gln Ala Gln
                                       40
Arg Asp Leu Tyr Arg Asp Val Met Leu Glu Asn Tyr Lys Asn Leu
                  50
                                      55
                                                           60
Ile Ile Leu Ala Gly Ser Glu Leu Phe Lys Arg Ser Leu Met Ser
                                                           75
                  65
                                       70
Gly Leu Glu Gln Met Glu Glu Leu Arg Thr Gly Val Thr Gly Val
                  80
                                      85
Leu Gln Glu Leu Asp Leu Gln Leu Lys Thr Lys Gly Ser Pro Leu
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95
                                     100
Leu Gln Asp Ile Ser Ala Glu Arg Ser Pro Asn Gly Val Gln Leu
                110
                                     115
Glu Arg Ser Asn Thr Ala Glu Lys Leu Tyr Asp Ser Asn His Ser
                125
                                     130
                                                          135
Gly Lys Val Phe Asn Glu His Pro Phe Leu Met Thr His Met Ile
                140
                                     145
Thr His Ile Gly Glu Lys Thr Ser Glu Asp Asn Gln Ser Gly
                155
                                     160
<210> 274
<211> 107
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<222> 5
<223> unknown or other
<400> 274
Met Trp Gln Val Xaa Ser Lys Ser Ser His Leu Ala Val His Gln
                                      10
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Arg Ile His Thr Gly Glu Lys Pro Tyr Lys Cys Asn Arg Cys Gly
                                      25
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Lys Cys Phe Ser Gln Ser Ser Leu Ala Arg His Gln Thr
                                                         Val
                 35
                                      40
                                                           45
His Thr Gly Glu Lys Pro Tyr Ile Cys Ala Glu Cys Gly Lys Ala
                 50
                                      55
                                                           60
Phe Ser Gln Lys Ser Asp Leu Val Val His Gln Ile Ile His Thr
                                                           75
                 65
                                      70
Gly Glu Lys Pro Asp Arg Cys Thr Val Cys Gly Lys Ala Phe
                                                         Ile
                 80
                                      85
Gln Lys Ser Gln Leu Thr Val His Gln Arg Ile His Thr Leu Met
                 95
                                     100
                                                          105
Lys Ser
<210> 275
<211> 105
<212> PRT
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<223> Incyte ID No: LI:789808.1.orf3:2000FEB01
<400> 275
Glu His Thr Asp Gly Lys Ser Tyr Ala Cys Ile Gln Cys Gly Lys
                    Tyr Ser Phe Thr Glu His Leu Arg Arg His
                Tyr
                                      25
Thr Gly Glu Lys Pro Phe Gly Cys Asn Glu Cys Gly Lys Thr Phe
                 35
                                      40
                                                           45
His Gln Lys Leu Ala Leu Ile Val His Gln Arg Thr His Ile Arg
                 50
                                      55
                                                           60
Gln Lys Pro Tyr Gly Cys Asn Glu Cys Gly Lys Ser Phe Cys Val
                                                           75
                                      70
                 65
Lys Ser Lys Leu Ile Ala His His Arg Thr Tyr Thr Gly Glu Lys
                 80
                                      85
Pro Tyr Glu Cys Asn Val Cys Gly Lys Leu Leu Ser Gln Asn
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<212> PRT
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<223> Incyte ID No: LI:792919.1.orf1:2000FEB01
<400> 276
His Gln Met Ile His Met Gly Gln Asn Pro Tyr Asn Cys Lys Glu
                                      10
                                                           15
Cys Gly Lys Ser Phe Lys Trp Ser Ser Tyr Leu Leu Val His Gln
                                      25
                                                           30
                 20
Arg Val His Thr Gly Glu Lys Pro Tyr Lys Cys Glu Glu Cys Gly
                                      40
                                                           45
                 35
Lys Gly Tyr Ile Ser Lys Ser Gly Leu Asp Phe His His Arg Thr
                 50
                                      55
His Thr Gly Glu Arg Ser Tyr Asn Cys Asp Asn Cys Gly Lys
                                                          Ser
                                      70
                                                           75
                 65
Phe Arg His Ala Ser Ser Ile Leu Asn His Lys Lys Leu His Cys
                                      85
                                                           90
                 80
Gln Arg Lys Pro Leu Lys Cys Glu Asp Cys Gly Lys Arg Leu Val
                                                          105
                 95
                                     100
Cys Arg Ser Tyr Cys Lys Asp Gln Gln Arg Asp His Ser Gly
                                                         Glu
                                                          120
                110
                                     115
Asn Pro Ser Lys Cys Glu Asp Cys Gly Lys Arg Tyr Lys Arg Arg
                125
                                     130
Leu Asn Leu Asp Ile Ile Leu Ser Leu Phe Leu Asn Asp Ile
                140
<210> 277
<211> 101
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Asp Thr Gly Thr Ser Trp Lys Pro Lys Met Gly Pro Leu Gln Phe
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Arg Asp Val Ala Ile Asp Phe Ser Gln Glu Glu Trp His Cys Leu
                 20
                                      25
                                                           30
Asp Thr Ala Gln Arg Asp Leu Tyr Arg Cys Val Met Leu Glu Asn
                                      40
                                                           45
Tyr Ser Asn Leu Val Phe Leu Gly Ile Thr Val Ser Lys Pro Asp
                 50
                                      55
                                                           60
Val Ile Ser Ser Leu Glu Gln Gly Arg Lys Pro Leu Thr Met Lys
                                                           75
                 65
                                      70
Arg Asn Glu Met Ile Ala Lys Pro Ser Val Ser Phe Leu Gln Val
                                      85
                 80
His Ser Glu Ser Gln Ser Pro Leu His Asp Ile
                 95
<210> 278
<211> 137
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<223> Incyte ID No: LI:794389.1.orf3:2000FEB01
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<222> 23
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Gly Leu Gln Lys Thr Phe Cys Arg Val Met Gln Phe Thr Leu His
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Arg Arg Ile His Thr Gly Glu Xaa Pro Tyr Glu Cys Lys Glu Cys
                 20
                                      25
                                                           30
Gly Lys Ser Phe Ser Ala His Ser Ser Leu Val Thr His Lys Arg
                                                           45
                 35
                                      40
Thr His Ser Gly Glu Lys Pro Tyr Lys Cys Lys Glu Cys Gly Lys
                                      55
                 50
                                                           60
Ala Phe Ser Ala His Ser Ser Leu Val Thr His Lys Arg Thr His
                                      70
                                                           75
                 65
Ser Gly Glu Lys Pro Tyr Thr Cys His Ala Cys Gly Lys Ala Phe
                                      85
                                                           90
                 80
Asn Thr Ser Ser Thr Leu Cys Gln His Asn Arg Ile His Thr Gly
                 95
                                     100
                                                          105
Glu Lys Pro Phe Gln Cys Ser Gln Cys Gly Lys Ser Phe Ser Cys
                                     115
                                                          120
                110
Ser Ser His Leu Thr Arg His Cys Arg Met Cys Asn Gly Lys Phe
                125
                                     130
                                                          135
Ser Lys
<210> 279
<211> 97
<212> PRT
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<220>
<221> unsure
<222> 4, 18, 23
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Leu Cys Ile Xaa Lys His Thr Gly Glu Lys Pro Tyr Glu Cys Tyr
                                      10
Ala Cys Xaa Asn Thr Phe Leu Xaa Lys Ser Asp Leu Ile Lys His
                                      25
                                                           3.0
Gln Arg Ile His Thr Gly Glu Lys Pro Tyr Glu Cys Asn Glu Cys
                                                           45
                 35
                                      40
Gly Lys Ser Phe Ser Glu Lys Ser Thr Leu Thr Lys His Leu Arg
                                      55
                 50
Thr His Arg Trp Glu Ile Leu Cys Met Tyr Ser Met Trp Lys
                                                          Ile
                                      70
                                                           75
                 65
Phe Leu Leu Leu Gln Phe His Arg Thr Ser Glu Lys Thr His
                                      85
                                                           90
                 80
Arg Gly Glu Thr Phe Trp Met
                 95
<210> 280
<211> 97
<212> PRT
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<220>
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<222> 93
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PCT/US01/06059

WO 01/62927 <223> unknown or other <400> 280 Leu Cys Ile Arg Lys His Thr Gly Glu Lys Pro Tyr Glu Cys Tyr 10 Ala Cys Gly Asn Thr Phe Leu Arg Lys Ser Asp Leu Ile Lys His 25 30 20 Gln Arg Ile His Thr Gly Glu Lys Pro Tyr Glu Cys Asn Glu Cys 35 40 Gly Lys Ser Phe Ser Glu Lys Ser Thr Leu Thr Lys His Leu Arg 50 55 60 Thr His Arg Trp Glu Ile Leu Cys Met Tyr Ser Met Trp Lys Ile 65 70 Phe Leu Leu Leu Gln Phe His Arg Thr Ser Glu Lys Thr His 80 85 Arg Gly Xaa Thr Phe Trp Met 95 <210> 281 <211> 179 <212> PRT <213> Homo sapiens <220> <221> misc\_feature <223> Incyte ID No: LI:796373.1.orf1:2000FEB01 Met Trp Glu Ser Phe Ser Gln Lys Thr Cys Leu Ile Ser His Gln 10 15 Arg Phe His Thr Gly Lys Thr Pro Phe Val Cys Thr Glu Cys Gly 20 25 30 Lys Ser Cys Ser His Lys Ser Gly Leu Ile Asn His Gln Arg Ile 45 35 40 His Thr Gly Glu Lys Pro Tyr Thr Cys Ser Asp Cys Gly Lys Ala 55 50 Phe Arg Asp Lys Ser Cys Leu Asn Arg His Arg Arg Thr His Thr 70 65 Gly Glu Arg Pro Tyr Gly Cys Ser Asp Cys Gly Lys Ala Phe Ser 80 85 90 His Leu Ser Cys Leu Val Tyr His Lys Gly Met Leu His Ala Arg 100 105 95 Glu Lys Cys Val Gly Ser Val Lys Leu Glu Asn Pro Cys Ser Glu 120 110 115 Ser His Ser Leu Ser His Thr Arg Asp Leu Ile Gln Asp Lys Asp 130 125 Ser Val Asn Met Val Thr Leu Gln Met Pro Ser Val Ala Ala Gln 140 145 Thr Ser Leu Thr Asn Ser Ala Phe Gln Ala Glu Ser Lys Val Ala 155 160 Ile Val Ser Gln Pro Val Ala Arg Ser Ser Val Ser Ala Asp 170 <210> 282 <211> 87 <212> PRT <213> Homo sapiens <220>

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<400> 282

Lys His Glu Ile Ile His Phe Glu Glu Glu Pro Ser Glu Tyr Asn 5 10 Asn Asn Gly Asn Ser Phe Trp Leu Asn Glu Asp Leu Ile Trp His

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Gln Lys Ile Lys Asn Trp Glu Gln Pro Phe Glu Tyr Asn Glu Cys
                 35
Gly Lys Ala Phe Pro Glu Asn Ser Leu Phe Leu Val His Lys Arg
                 50
                                      55
                                                           60
Ala Tyr Thr Gly Gln Lys Thr Cys Lys Tyr
                                         Thr Glu His Gly Lys
                                      70
                 65
Thr Cys Tyr Met Ser Phe Phe Ile Thr His Gln Gln
                                      85
                 80
<210> 283
<211> 172
<212> PRT
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Asn Glu Cys Gly Lys Ala Leu Ser Ser His Ser Thr Leu Ile Ile
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His Glu Arg Ile His Thr Gly Glu Lys Pro Cys Lys Cys Lys Val
                                       25
                 20
Cys Gly Lys Ala Phe Arg Gln Ser Ser Ala Leu Ile Gln His Gln
                                       40
                                                           45
                 35
Arg Met His Thr Gly Glu Arg Pro Tyr Lys Cys Asn Glu Cys Asp
                 50
                                       55
                                                           60
Lys Thr Phe Arg Cys Asn Ser Ser Leu Ser Asn His Gln Arg Ile
                                                           75
                                      70
                  65
His Thr Gly Glu Lys Pro Tyr Arg Cys Leu Glu Cys Gly Met Ser
                                                           90
                 80
                                      85
Phe Gly Gln Ser Ala Ala Leu Ile Gln His Gln Arg Ile His Three
                                                          105
                 95
                                      100
Gly Glu Lys Pro Phe Lys Cys Asn Thr Cys Gly Lys Thr Phe Arg
                110
                                     115
                                                          120
Gln Ser Ser Leu Ile Ala His Gln Arg Ile His Thr Gly Glu
                                                          135
                 125
                                      130
Lys Pro Tyr Glu Cys Asn Ala Cys Gly Lys Leu Phe Ser Gln Arg
                                                          150
                                      145
                 140
Ser Ser Leu Thr Asn His Tyr Lys Ile His Ile Glu Glu Asp Ser
                 155
                                                          165
                                     160
Leu Lys Ala Asp Leu His Val
                 170
<210> 284
<211> 151
<212> PRT
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<220>
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<222> 104
<223> unknown or other
<400> 284
Lys Ile Met His Thr Gly Glu Lys Arg Tyr Glu Cys Asp Asp Cys
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Gly Gly Thr Phe Arg Ser Ser Ser Ser Leu Arg Val His Lys Arg
                                       25
                  20
Ile His Thr Gly Glu Lys Pro Tyr Lys Cys Glu Glu Cys
                                                      Gly Lys
                                       40
                  35
Ala Tyr Met Ser Tyr Ser Ser Leu Ile Asn His Lys Ser Thr His
                                       55
                                                            60
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Ser Gly Glu Lys Asn Cys Lys Cys Asp Glu Cys Gly Lys Ser Phe
                                                          75
                 65
Asn Tyr Ser Ser Val Leu Asp Gln His Lys Arg Ile His Thr Gly
                 80
                                      85
                                                           90
Glu Lys Pro Tyr Glu Cys Gly Glu Cys Gly Lys Ala Phe Xaa Asn
                 95
                                     100
Ser Ser Gly Leu Arg Val His Lys Arg Ile His Thr Gly Glu Lys
                110
                                     115
                                                          120
Pro Tyr Glu Cys Asp Ile Cys Gly Lys Thr Phe Ser Asn Ser Ser
                125
                                     130
                                                          135
Gly Leu Thr Val His Lys Arg Ile His Thr Val Ser Asp Glu Leu
                140
                                     145
                                                          150
Pro
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<211> 89
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Leu Thr Tyr Leu Arg Lys Lys Leu Arg Gly Arg Gly Lys Lys Glu
Glu Glu Gly Met Ala Leu Ser Gln Gly Leu Phe Thr Phe Lys Asp
                                      25
                                                           30
Val Ala Ile Glu Phe Ser Gln Glu Glu Trp Glu Cys Leu Asp Pro
                 35
                                      40
                                                           45
Ala Gln Arg Ala Leu Tyr Arg Asp Val Met Leu Glu Asn Tyr Arg
                                      55
                                                           60
                 50
Asn Leu Leu Ser Leu Asp Glu Asp Asn Ile Pro Pro Glu Asp
                 65
                                      70
Ser His Leu Ala Ala Cys Gly Gln Ser Thr Leu Pro Leu Pro
                 80
<210> 286
<211> 146
<212> PRT
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<223> Incyte ID No: LI:800894.1.orf2:2000FEB01
<400> 286
Pro Ala Gly Ile Gly Arg Ser Thr Thr Lys Ser Pro Gly Pro Pro
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Gly Ser Leu Glu Met Gly Ser Leu Thr Phe Arg Asp Val Ala Ile
                                      25
Glu Phe Ser Leu Glu Glu Trp Gln Cys Leu Asp Thr Ala Gln Gln
                                      40
                                                           45
                 35
Asn Leu Tyr Arg Asn Val Met Leu Glu Asn Tyr Arg Asn Leu Val
                  50
                                      55
                                                           60
Phe Leu Gly Ile Ala Ala Phe Lys Pro Asp
                                         Leu Ile Ile Phe Leu
                  65
                                      70
                                                           75
Glu Glu Gly Lys Glu Ser Trp Asn Met Lys Arg His Glu Met Val
                                      85
                                                           90
Glu Glu Ser Pro Val Ile Cys Ser His Phe Ala Gln Asp Leu Trp
                  95
                                     100
                                                          105
Pro Glu Gln Gly Ile Glu Asp Ser Phe Gln Lys Val Ile Leu Arg
                 110
                                     115
Arg Tyr Lys Ile His His His Ala Cys Glu Leu Gly Pro Ile Met
                 125
                                     130
Asn His Tyr Pro Thr Cys Gly Gln Met His Ile
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140 145

<210> 287 <211> 78 <212> PRT . <213> Homo sapiens <220> <221> misc\_feature <223> Incyte ID No: LI:801015.1.orf1:2000FEB01 <400> 287 Gly Ser Arg Lys Met Asp Ser Val Ala Phe Glu Asp Val Ala Val 10 15 Asn Phe Thr Gln Glu Glu Trp Ala Leu Leu Asp Pro Trp Gln Lys 25 30 20 Lys Leu Tyr Arg Asp Val Met Leu Glu Thr Tyr Arg Asn Leu Ala 35 40 45 Ser Val Gly Asp Asp Asp Asn Ile Pro Ser Leu Arg Glu Gln Val 55 60 50 Ala His Gln Arg Tyr Phe Lys Thr Trp His Val Glu Arg Glu Tyr 65 Phe Ser Lys <210> 288 <211> 126 <212> PRT <213> Homo sapiens <221> misc\_feature <223> Incyte ID No: LI:801236.1.orf3:2000FEB01 <220> <221> unsure <222> 4 <223> unknown or other <400> 288 Met Trp Glu Xaa Phe Ser His Thr Pro Ala Phe Ile Gln His Gln 15 10 Arg Ile His Thr Gly Glu Lys Pro Tyr Glu Cys Asn Ala Cys Gly 25 20 Lys Ala Phe Asn Arg Ser Ala His Leu Thr Glu His Gln Arg Thr 40 His Thr Gly Glu Lys Pro Tyr Val Cys Lys Glu Cys Gly Lys Thr 55 60 Phe Ser Arg Ser Thr His Leu Thr Glu His Leu Lys Ile His Ser 75 65 70 Cys Val Lys Pro Tyr Gln Cys Asn Glu Cys Gln Lys Leu Phe Cys 90 85 80 Tyr Arg Thr Ser Leu Ile Arg His Gln Arg Thr His Thr Gly Glu 100 105 95 Lys Pro Tyr Gln Cys Asn Glu Cys Gly Lys Ser Phe Ser Leu Ser 110 Ser Ala Leu Thr Lys His 125 <210> 289 <211> 96 <212> PRT <213> Homo sapiens <220> <221> misc\_feature <223> Incyte ID No: LI:803335.1.orf1:2000FEB01

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 Ser Ala Ala Met Phe Pro Val Phe Ser Gly Cys Phe Gln Glu Leu
Gln Glu Lys Asn Lys Ser Leu Glu Leu Val Ser Phe Glu Glu Val
                  20
                                       25
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Ala Val His Phe Thr Trp Glu Glu Trp Gln Asp Leu Asp Asp Ala
                                       40
                  35
                                                            45
Gln Arg Thr Leu Tyr Arg Asp Val Met Leu Glu Thr Tyr Ser Ser
                  50
                                       55
                                                            60
 Leu Val Ser Leu Gly His Cys Ile Thr Lys Pro Glu Met Ile Phe
                  65
                                       70
                                                            75
Lys Leu Glu Gln Gly Ala Glu Pro Trp Ile Val Glu Glu Thr Leu
                  80
Asn Leu Arg Leu Ser Gly
                  95
 <210> 290
<211> 149
<212> PRT
 <213> Homo sapiens
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Lys Asn Ser Tyr Trp Arg Lys Asn Pro Thr Asn Met Lys Asn Val
                                       10
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 Ala Lys Leu Leu Ile Asn Ser Gln Arg Leu Leu Asn Ile Arg Glu
                                       25
                  20
 Phe Val Gln Glu Gly Asn Pro Thr Asn Leu Lys Asn Val Ala Ser
                                       40
Leu Leu Ala Ile Pro Gln Ser Leu Leu Asn Ile His Val Ile His
                  50
                                       55
                                                            60
 Thr Gly Gly Asn Ser Tyr Asn Cys Val Glu Cys Cys Asn Ala Leu
                  65
                                       70
                                                            75
Asn Gln Ser Leu Arg Leu Thr Thr Tyr Lys Thr Thr His Thr Gly
                                                            90
                  80
                                       85
 Glu Lys Pro Cys Met Cys Glu Glu Cys Gly Lys Ala Ser Asn Arg
                  95
                                      100
                                                           105
Ser Ser Ile Leu Lys Arg His Lys Leu Ile His Thr Gln Glu Arg
                 110
                                      115
                                                           120
 Leu Tyr Lys Pro Glu Arg Cys Asp Asn Ala Phe Gly Asn Thr Ser
                 125
                                      130
 Asp Phe Ser Glu Tyr Lys Arg Asn Arg Thr Asp Glu Lys Ser
                 140
                                      145
 <210> 291
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 Trp Trp Glu Ile Cys Ala His Ser Asp Val Ala Ala Glu Glu Gly
                                       10
 Lys Ala Arg Arg Ser Arg Gln His Arg Phe Leu Gly Thr Cys Glu
                  20
                                       25
                                                            30
Gly Ile Met Arg Arg Ala Glu Leu Ser Ser Gln Val Glu Asp Ser
                                       40
                                                            45
 Thr Leu His Ala Trp Ile Arg Tyr Ser Leu Val Leu Asp Val Asp
                  50
                                       55
                                                            60
Cys Trp His Ile Ala Ala Gln Leu Glu Met Tyr Gly Cys Pro His
                  65
                                       70
```

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Leu Asp Leu Thr Glu Ser Arg Gly Ala Ala Arg Lys Leu His
Leu Leu Gly Phe Ser Ala Leu Pro Thr Leu Val Asp Met Ile Thr
                 95
                                     100
                                                          105
Ser Gln Gly Ser Val Ser Phe Arg Asp Val Thr Met Gly Phe Thr
                110
                                     115
                                                          120
Gln Glu Glu Trp His His Leu Asp Pro Ala Gln Arg Thr Leu
                125
<210> 292
<211> 212
<212> PRT
<213> Homo sapiens
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His Asn Phe Gln Leu Gln Lys His His Arg Ile His Thr Gly Glu
                                      10
Lys Pro Phe Lys Cys Glu Ile Cys Gly Lys Ser Phe Cys Leu Arg
                 20
                                                           3.0
Ser Ser Leu Asn Arg His Cys Met Val His Thr Ala Glu Lys Leu
                 35
                                      40
                                                           45
Tyr Lys Ser Glu Lys Tyr Gly Arg Gly Phe Ile Asp Arg Leu Asp
                                                           60
                 50
                                      55
Leu His Lys His Gln Met Ile His Met Gly Gln Lys Pro Tyr Asn
                                      70
                                                           75
                 65
Cys Lys Glu Cys Gly Lys Ser Phe Lys Trp
                                        Ser Ser Tyr Leu Leu
                                      85
                                                           90
                 80
Val His Gln Arg Val His Thr Gly Glu Lys Pro Tyr Lys Cys Glu
                                     100
                                                          105
Glu Cys Gly Lys Gly
                    Tyr Ile Ser Lys Ser Gly Leu Asp Phe His
                110
                                     115
                                                          120
His Arg Thr His Thr Gly Glu Arg Ser Tyr Asn Cys Asp Asn Cys
                                     130
                                                          135
                125
Gly Lys Ser Phe Arg His Ala Ser Ser Ile Leu Asn His
                                                     Lys Lys
                                     145
                                                          150
                140
Leu His Cys Gln Arg Lys Pro Leu Lys Cys Glu Asp Cys Gly Lys
                155
                                     160
                                                          165
Arg Leu Val Cys Arg Ser Tyr Cys Lys Asp Gln Gln Arg Asp His
                170
                                     175
                                                          180
Ser Gly Glu Asn Pro Ser Lys Cys Glu Asp Cys Gly Lys Arg
                                                          Tyr
                185
                                     190
                                                          195
Lys Arg Arg Leu Asn Leu Asp Ile Ile Leu Ser Leu Phe Leu Asn
                                     205
                                                          210
                200
Asp Ile
<210> 293
<211> 108
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Lys Pro Tyr Met Cys Lys Glu Cys Arg Lys Thr Phe Ser Gln Asn
 1
                                      10
                                                           15
Ala Gly Leu Ala Gln His Gln Arg Ile His Thr Gly Glu Lys Pro
                                                           30
                 20
                                      25
Tyr Glu Cys Asn Val Cys Gly Lys Ala Phe Ser Tyr Ser Gly Ser
                 35
                                      40
Leu Thr Leu His Gln Arg Ile His Thr Gly Glu Arg Pro Tyr Glu
```

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50
                                                           60
Cys Lys Asp Cys Arg Lys Ser Phe Arg Gln Arg Ala His Leu Ala
                 65
                                      70
                                                           75
His His Glu Arg Ile His Thr Met Glu Ser Phe Leu Thr Leu Ser
                 80
                                      85
                                                           90
Ser Pro Ser Pro Ser Thr Ser Asn Gln Leu Pro Arg Pro Val Gly
                                     100
                                                          105
                 95
Phe Ile Ser
<210> 294
<211> 83
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Pro Ala Cys Thr Gly Gly Phe Ala Gly Arg Met Ser Gly His Pro
Gly Ser Trp Glu Met Asn Ser Val Ala Phe Glu Asp Val Ala Val
                                      25
                                                           30
Asn Phe Thr Gln Glu Glu Trp Ala Leu Leu Asp Pro Ser Gln Lys
                                      40
                                                           45
Asn Leu Tyr Arg Asp Val Met Gln Glu Thr Phe Arg Asn Leu Ala
                                      55
                                                           60
                 50
Ser Ile Gly Asn Lys Gly Glu Asp Gln Ser Ile Glu Asp Gln
                 65
Lys Asn Ser Ser Arg Asn Leu Arg
                 80
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Pro Tyr Val Cys Lys Glu Cys Gly Lys Ala Phe Thr Gln Tyr Ser
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Gly Leu Ser Met His Val Arg Ser His Ser Gly Asp Lys Pro Tyr
                 20
                                      25
                                                           3.0
Glu Cys Lys Glu Cys Gly Lys Ser Phe Leu Thr Ser Ser Arg Leu
                  35
                                      40
                                                           45
Ile Gln His Ile Arg
                    Thr His Thr Gly Glu Lys Pro Phe Val Cys
                  50
                                      55
                                                           60
Val Glu Cys Gly Lys Ala Phe Ala Val Ser Ser Asn Leu Ser Gly
                                      70
                                                           75
                  65
His Leu Arg Thr His Thr Glu Glu Lys Ala Cys Glu Cys Lys Ile
                                      85
                 80
Cys Gly Lys Val Phe Gly Tyr Pro Ser Cys Leu Asn Asn His Met
                                     100
                  95
Arg Thr His Ser Ala Gln Lys Pro Tyr Thr Cys Lys Glu Cys Gly
                                                          120
                 110
                                     115
                     Ser Thr His Leu Lys Ile His Met Arg Ile
Lys Ala Phe Asn Tyr
                125
                                     130
                                                          135
His Thr Gly Glu Lys Pro Tyr Glu Cys Lys Gln Cys Gly Lys Ala
                 140
                                     145
                                                          150
Phe Ser His Ser Ser Ser Phe Gln Ile His Glu Arg Thr His
                155
                                     160
                                                          165
                     Glu Cys Lys Glu Cys Gly Lys Ala Phe Thr
Gly Glu Lys Pro Tyr
                170
                                     175
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<211> 97
<212> PRT
<213> Homo sapiens
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Leu Cys Ile Ser Lys His Pro Gly Glu Lys Pro Tyr Glu Cys Tyr
                                      10
Ala Cys Gly Asn Thr Phe Leu Arg Lys Ser Asp Leu Ile Lys His
                                      25
                                                           30
                 20
Gln Arg Ile His Thr Gly Glu Lys Pro Tyr Glu Cys Asn Glu Cys
                 35
                                      40
                                                           45
Gly Lys Ser Phe Ser Glu Lys Ser Thr Leu Thr Lys His Leu Arg
                 50
                                      55
                                                           60
Thr His Arg Trp Glu Ile Leu Cys Met Tyr
                                        Ser Met Trp Lys
                                      70
                                                           75
                 65
Phe Leu Leu Leu Gln Phe His Arg Thr Ser Glu Lys Pro His
                 80
                                      85
Arg Gly Glu Thr Phe Trp Met
<210> 297
<211> 217
<212> PRT
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Ala Ala Pro Glu Ser Gln Gln Gln Arg Asn Arg Arg Gly Glu Arg
                                      10
Pro Phe Thr Cys Met Glu Cys Gly Lys Ser Phe Arg Leu Lys Ile
                                      25
                 20
Asn Leu Ile Ile His Gln Arg Asn His Ile Lys Glu Gly Pro Tyr
                                      40
Glu Cys Ala Glu Cys Glu Ile Ser Phe Arg His Lys Gln Gln Leu
                 50
                                      55
                                                           60
Thr Leu His Gln Arg Ile His Arg Val Arg Gly Gly Cys Val Ser
                                      70
                 65
                                                           75
Pro Glu Arg Gly Pro Thr Phe Asn Pro Lys His Ala Leu Lys Pro
                                      85
                                                           90
                 80
Arg Pro Lys Ser Pro Ser Ser Gly Ser Gly Gly Gly Pro Lys
                 95
                                     100
                                                          105
Pro Tyr Lys Cys Pro Glu Cys Asp Ser Ser Phe Ser His Lys Ser
                                                          120
                110
                                     115
Ser Leu Thr Lys His Gln Ile Thr His Thr Gly Glu Arg Pro Tyr
                125
                                     130
                                                          135
Thr Cys Pro Glu Cys Lys Lys Ser Phe Arg Leu His Ile Ser Leu
                140
                                     145
                                                          150
Val Ile His Gln Arg Val His Ala Gly Lys His Glu Val Ser Phe
                                                          165
                155
                                     160
Ile Cys Ser Leu Cys Gly Lys Ser Phe Ser Arg Pro Ser His Leu
                170
                                     175
Leu Arg His Gln Arg Thr His Thr Gly Glu Arg Pro Phe Lys Cys
                                                          195
                185
                                     190
Pro Glu Cys Glu Lys Ser Phe Ser Glu Lys Ser Lys Leu Thr Asn
                                     205
                                                          210
                200
His Cys Arg Val His Ser Arg
                215
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<211> 137
<212> PRT
<213> Homo sapiens
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<223> Incyte ID No: LG:892274.1.orf3:2000MAY19
<400> 298
Asp Gly Gly Leu Asp Leu Gly Pro Thr Asn Ser Glu Gly Ile Pro
Ser Pro Asp Leu Asn Pro Val Leu Gly Met Gly Ser Trp Arg His
                                      25
                                                           30
Ile Asp Ser Ile Thr Pro Gly Thr Pro Gly Ser Ala Gly Leu Asp
                                      40
                                                           45
                 35
Leu Pro Ala Arg Glu Arg Ile Thr Leu Val Gly Gly Asp Lys Pro
                 50
                                      55
                                                           60
Ile Lys Val Pro Thr Gly Ile Trp Gly Thr Ser Pro Ala Gly Tyr
                 65
                                      70
Met Gly Leu Ile Leu Gly Lys Ser Arg Leu Asn Leu Gln Gly Met
                                                           90
                 80
                                      85
Thr Val Val Pro Gly Ala Val Asp Ser Asp Tyr Glu Gly Glu Thr
                                                          105
                 95
                                     100
Gln Val Val Leu Met Ser Gln Asp Leu Trp Val Phe Glu Leu Gly
                110
                                     115
                                                          120
Glu Tyr Ile Ala Gln Leu Leu Ile Pro Cys Lys Leu His Pro
                                     130
                                                          135
Ser Pro
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Pro Lys Gln Gly Ile Asn Val Trp Ser Pro Arg His Pro Glu Asn
                                      10
                                                           15
Phe Leu Gly Ile Glu Ser Arg Pro Pro Met Leu Ser Leu Ser Pro
                  20
                                      25
                                                           3.0
Ile Leu Leu Tyr Thr Cys Glu Met Phe Gln Asp Pro Val Ala Phe
                  35
                                      40
Lys Asp Val Ala Val Asn Phe Thr Gln Glu Glu Trp Ala Leu Leu
                                      55
                                                           60
                  50
Asp Ile Ser Gln Lys Asn Leu Tyr Arg Glu Val Met Leu Glu Thr
                                                           75
                  65
                                      70
Phe Trp Asn Leu Thr Ser Ile Gly Lys Lys Trp Lys Asp Gln Asn
                                      85
                 80
Ile Glu Tyr Glu Tyr Gln Asn Pro Arg Arg Asn Phe Arg Ser Val
                                     100
                                                          105
Thr Glu Glu Lys Val Asn Glu Ile Lys Glu Asp Ser His Cys Gly
                 110
                                     115
                                                          120
Glu Thr Phe Thr Pro Val Pro Asp Asp Arg Leu Asn Phe Gln Lys
                 125
                                     130
                                                          135
Lys Lys Ala Ser Pro Glu Val Lys Ser Cys Asp Ser Phe Val Cys
                                                          150
                 140
                                     145
Glu Val Gly Leu Gly Asn Ser Ser Ser Asn Met Asn Ile Arg Gly
                                     160
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Asp Thr Gly His
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<210> 300

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Asp Ala Trp Ala Arg Pro Pro Val Leu Ser Leu Ser Pro Ile Leu
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Leu Tyr Thr Cys Glu Met Phe Gln Asp Pro Val Ala Phe Asp Asp
                                      25
                 20
Val Ala Val Asn Phe Thr Gln Glu Glu Trp Ala Leu Leu Asp Ile
                                      40
                 35
                                                           45
Ser Gln Arg Lys Leu Tyr Lys Glu Val Met Leu Glu Thr Phe Arg
                 50
                                      55
                                                           60
Asn Leu Thr Ser Val Gly Lys Ser Trp Lys Asp Gln Asn Ile Glu
                                      70
                 65
Tyr Glu Tyr Gln Asn Pro Arg Arg Asn Phe Arg Ser Leu Ile Glu
                                                           90
                 80
                                      85
Lys Lys Val Asn Glu Ile Lys Asp Asp Ser His Cys Gly Glu Thr
                 95
                                     100
                                                          105
Phe Thr Gln Val Pro Asp Asp Arg Leu Asn Phe Gln Glu Lys Lys
                                                          120
                110
                                     115
Ala Ser Pro Glu Ile Lys Ser Cys Asp Ser Phe Val Cys Gly Lys
                125
<210> 301
<211> 170
<212> PRT
<213> Homo sapiens
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Thr Val Met Leu Cys Asp Glu Glu Ala Gln Lys Arg Lys Ala Lys
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                                      10
                                                           15
Glu Ser Gly Met Ala Leu Pro Gln Gly Arg Leu Thr Phe Met Asp
                                                           30
                 20
                                      25
Val Ala Ile Glu Phe Ser Gln Glu Glu Trp Lys Ser Leu Asp Pro
                                      40
                                                           45
                 35
Gly Gln Arg Ala Leu Tyr Arg Asp Val Met Leu Glu Asn Tyr Arg
                 50
                                      55
                                                           60
Asn Leu Val Phe Leu Gly Ile Cys Leu Pro Asp Leu Ser Ile Ile
                                      70
                                                           75
                 65
Ser Met Leu Lys Gln Arg Arg Glu Pro Leu Ile Leu Gln Ser Gln
                 80
                                      85
                                                           90
Val Lys Ile Val Lys Asn Thr Asp Gly Arg Glu Cys Val Arg Ser
                                     100
                 95
Val Asn Thr Gly Arg Ser Cys Val Leu Gly Ser Asn Ala Glu Asn
                                                          120
                                     115
                110
Lys Pro Ile Lys Asn Gln Leu Gly Leu Thr Leu Glu Ser His Leu
                                     130
                125
Ser Glu Leu Gln Leu Phe Gln Ala Gly Arg Lys Ile Tyr Arg Ser
                                                          150
                140
                                     145
Asn Pro Val Glu Lys Phe Thr Asn His Arg Ser Ser Val Ser Pro
                155
                                     160
                                                          165
Leu Gln Lys Ile Ser
                170
<210> 302
<211> 181
<212> PRT
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Thr Ser Tyr Ile Arg Thr Lys Thr Tyr Glu Cys Asn Ile Cys Glu
                                                          15
                                      10
Lys Ile Phe Lys Gln Pro Ile His Leu Thr Glu His Met Arg Ile
                                                          30
                 20
                                      25
His Thr Gly Glu Lys Pro Phe Arg Cys Lys Glu Cys Gly Arg Ala
                 35
                                      40
Phe Ser Gln Ser Ala Ser Leu Ser Thr His Gln Arg Ile His Thr
                 50
                                      55
Gly Glu Lys Pro Phe Glu Cys Glu Cys Gly Lys Ala Phe Arg
                 65
His Arg Ser Ser Leu Asn Gln His His Arg Thr His Thr Gly Glu
                                                          90
                                      85
Lys Pro Tyr Val Cys Asp Lys Cys Gln Lys Ala Phe Ser Gln Asn
                 95
                                     100
                                                          105
Ile Ser Leu Val Gln His Leu Arg Thr His Ser Gly Glu Lys Pro
                110
                                     115
                                                          120
Phe Thr Cys Asn Glu Cys Gly Lys Thr Phe Arg Gln Ile Arg His
                125
                                     130
                                                          135
Leu Ser Glu His Ile Arg Ile His Thr Gly Glu Lys Pro Tyr Ala
                                     145
                140
Cys Thr Ala Cys Cys Lys Thr Phe Ser His Arg Ala Tyr Leu Thr
                155
                                     160
                                                          165
His His Gln Arg Ser Ile Leu Gly Arg Asp Leu Gln Cys Lys Glu
                170
Cys
<210> 303
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Ala Phe Ser Arg Cys Ser Ser Leu Val Gln His Glu Arg Thr His
Thr Gly Glu Lys Pro Phe Glu Cys Ser Ile Cys Gly Arg Ala Phe
                 20
                                      25
                                                           30
Gly Gln Ser Pro Ser
                    Leu Tyr Lys His Met Arg Ile His Lys Arg
                 35
                                      40
Gly Lys Pro Tyr Gln Ser Ser Asn Tyr Ser Ile Asp Phe Lys His
                                                           60
                 50
                                      55
Ser Thr Ser Leu Thr Gln Asp Glu Ser Thr Leu Thr Glu Val Lys
                 65
                                      70
Ser Tyr His Cys Asn Asp Cys Gly Glu Asp Phe Ser His Ile Thr
                                      85
                                                           90
                 80
Asp Phe Thr Asp His Gln Arg Ile His Thr Ala Glu Asn Pro Tyr
                 95
                                     100
                                                          105
Asp Cys Glu Gln Ala Phe Ser Gln Gln Ala Ile Ser His Pro Gly
                110
                                     115
                                                          120
Glu Lys Pro Tyr Gln Cys Asn Val Cys Gly Lys Ala Phe Lys Arg
                125
                                     130
                                                          135
Ser Thr Ser Phe Ile Glu His His Arg Ile His Thr Gly Glu Lys
                140
                                     145
                                                          150
Pro Tyr Glu Cys Asn Glu Cys Gly Glu Ala Phe Ser Arg Arg Ser
                155
                                                          165
                                     160
Ser Leu Thr Gln His Glu Arg Thr His Thr Gly Glu Lys Pro Tyr
```

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170
                                                          180
                                     175
Glu Cys Ile Asp Cys Gly Lys Ala Phe Ser Gln Ser Ser Leu
                185
                                     190
                                                          195
Ile Gln His Glu Arg Thr His Thr Gly Glu Lys Pro Tyr Glu Cys
                200
                                     205
Asn Glu Cys Gly Arg Ala Phe Arg Lys Lys Thr Asn Leu His Asp
                215
                                     220
                                                          225
His Gln Arg Ile His Thr Gly Glu Lys Pro Tyr Ser Cys Lys
                                                          Glu
                230
                                     235
                                                          240
Cys Gly Lys Asn Phe Ser Arg Ser Ser Ala Leu Thr Lys His
                                                          Gln
                245
                                     250
                                                          255
Arg Ile His Thr Arg Asn Lys Leu
                260
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Arg Leu Leu Val Pro Glu Glu Glu Thr Gln Lys Arg Lys Arg Lys
Ala Lys Glu Ser Gly Met Ala Leu Ser Gln Gly Leu Leu Thr Phe
                                      25
                                                           30
                 20
Arg Asp Val Ala Ile Glu Phe Ser Glu Glu Glu Trp Lys Cys Leu
                 35
                                      40
                                                           45
Asp Pro Ala Gln Arg Thr Leu Tyr Arg Asp Val Met Leu Glu Asn
                 50
                                      55
                                                           60
Tyr Arg Asn Leu Val Ser Leu Asp Ile Ser Ser Lys Cys Thr Met
                 65
                                      70
Lys Glu Phe Leu Ser Thr Ala Gln Gly Asn Arg Glu Val Phe His
                                      85
                 80
                                                           90
Ala Gly Thr Leu Gln Ile His Glu Ser His His Asn Gly Asp Phe
                 95
                                     100
                                                          105
Cys Tyr Gln Asp Val Asp Lys Asp Ile His Asp Tyr Glu Phe Gln
                110
                                     115
                                                          120
Trp Gln Glu Asp Glu Arg Asn Gly His Glu Ala Pro Met Thr Lys
                125
                                     130
                                                          135
Ile Lys Lys Leu Thr Gly Ile Thr Glu Arg Tyr Asp Gln Ser His
                                     145
                                                          150
                140
Ala Arg Asn Lys Pro Ile Lys Asp Gln Leu Gly Ser Ser Phe His
                155
Ser His Leu Pro Glu Met His Ile Phe Gln Thr Glu Glu Lys Ile
                170
                                     175
                                                          180
Asp Asn Gln Val Val Lys Ser Ile His Asp Ala Ser Leu Val Ser
                                                          195
                185
                                     190
Thr Ala Gln Arg Ile Ser Cys Arg Pro Lys Thr His Ile Ser Asn
                                     205
                                                          210
                200
Asn His Gly Asn Asn Phe Trp Asn Ser Ser Leu Leu Thr Gln Lys
                                     220
                                                          225
                215
Gln Glu Val His Met Arg Glu Lys Ser Phe Gln Cys Asn Glu Ser
                230
                                      235
Gly Lys Ala Phe Asn Tyr Ser Ser Leu Leu Arg Lys His Gln Ile
                                                          255
                245
                                     250
Ile His Leu Ala Asp Lys Tyr Lys Cys Asp Val Cys Gly Lys Leu
                 260
                                      265
                                                           270
Phe Asn Gln Lys Arg Asn Leu Ala Cys His Arg Arg Cys His Thr
                275
                                     280
                                                          285
Gly Glu Asn Pro Tyr Lys Cys Asn Glu Cys Gly Lys Thr Phe Ser
                                                          300
                 290
                                     295
Gln Thr Ser Ser Leu Thr Cys His Arg Arg Leu His Thr Gly Glu
                305
                                     310
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Lys Pro Tyr Lys Cys Glu Glu Cys Asp Lys Ala Phe His Phe Lys

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320
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Ser Ile Leu Glu Arg His Arg Ile Ile His
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Leu Thr Tyr Leu Arg Lys Lys Leu Arg Gly Arg Gly Lys Lys Glu
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Glu Glu Gly Met Ala Leu Ser Gln Gly Leu Phe Thr Phe Lys Asp
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                                      25
Val Ala Ile Glu Phe Ser Gln Glu Glu Trp Glu Cys Leu Asp Pro
                 35
                                      40
Ala Gln Arg Ala Leu Tyr Arg Asp Val Met Leu Glu Asn Tyr Arg
                                      55
                                                           60
                 50
Asn Leu Leu Ser Leu Asp Glu Asp Asn Ile Pro Pro Glu Asp
                                                         Glv
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Ser His Leu Ala Ala Cys Gly Gln Ser Thr Leu Pro Leu Pro
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Gly Thr Ser Ala Val Phe Asn Pro Ser Val Leu His Tyr Gln Gln
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Ala Leu Thr Ser Ala Gln Leu Gln Gln His Ala Ala Phe Ile Pro
                 35
                                      40
                                                           45
Thr Gly Met Cys Pro Tyr Cys Pro Thr Ser Cys Ala Leu Leu Val
                                      55
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Met Cys Phe Leu Leu Ile Ser Leu Ser Cys Leu Val Ala Ser Ser
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Leu Leu Leu Lys Val
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<211> 386
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Ala Leu Asp Gly Val Arg Pro Pro Gln Ser Gln Glu Met Ala Thr
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Ala Val Glu Pro Glu Asp Gln Asp Leu Trp Glu Glu Glu Gly Ile
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                                                           30
Leu Met Val Lys Leu Glu Asp Asp Phe Thr Cys Arg Pro Glu Ser
                                      40
                 35
Val Leu Gln Arg Asp Asp Pro Val Leu Glu Thr Ser His Gln Asn
```

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50
Phe Arg Arg Phe Arg Tyr Gln Glu Ala Ala Ser Pro Arg Glu Ala
                 65
                                      70
Leu Ile Arg Leu Arg Glu Leu Cys His Gln Trp Leu Arg Pro Glu
                 80
                                      85
Arg Arg Thr Lys Glu Gln Ile Leu Glu Leu Leu Val Leu Glu Gln
                 95
                                     100
                                                          105
Phe Leu Thr Val Leu Pro Gly Glu Leu Gln Ser Trp Val Arg Gly
                110
                                     115
                                                          120
Gln Arg Pro Glu Ser Gly Glu Glu Ala Val Thr Leu Val Glu Gly
                                                          135
                125
                                     130
Leu Gln Lys Gln Pro Arg Arg Pro Arg Arg Trp Val Thr Val His
                                                          150
                                     145
                140
Val His Gly Gln Glu Val Leu Ser Glu Glu Thr Val His Leu Gly
                155
                                     160
                                                          165
Ala Glu Pro Glu Ser Pro Asn Glu Leu Gln Asp Pro Val Gln Ser
                                     175
                170
Ser Thr Pro Glu Gln Ser Pro Glu Glu Thr Thr Gln Ser Pro Asp
                                                          195
                185
                                     190
Leu Gly Ala Pro Ala Glu Gln Arg Pro His Gln Glu Glu Leu
                200
                                     205
                                                          210
Gln Thr Leu Gln Glu Ser Glu Val Pro Val Pro Glu Asp Pro Asp
                215
                                     220
                                                          225
Leu Pro Ala Glu Arg Ser Ser Gly Asp Ser Glu Met Val Ala Leu
                                                          240
                230
                                     235
Leu Thr Ala Leu Ser Gln Gly Leu Val Thr Phe Lys Asp Val Ala
                                     250
                                                          255
                245
Val Cys Phe Ser Gln Asp Gln Trp Ser Asp Leu Asp Pro Thr Gln
                                     265
                                                          270
                260
Lys Glu Phe Tyr Gly Glu Tyr Val Leu Glu Glu Arg Leu Trp Asn
                275
                                     280
                                                          285
Cys Cys Leu Ser Val His Ser Gln Ser Pro Arg Pro Asp Glu Ile
                290
                                     295
                                                          300
Leu Pro Gly Leu Asp Glu Glu Glu Pro Gly Val Pro Asp Ile Gln
                305
                                     310
                                                          315
Glu Pro Gln Glu Thr Gln Glu Pro Glu Ile Leu Ser Phe Thr
                                                          Tyr
                                     325
                320
Thr Gly Asp Arg Ser Lys Asp Glu Glu Glu Cys Leu Glu Gln Glu
                335
                                     340
Asp Leu Ser Leu Glu Asp Ile His Arg Pro Val Leu Gly Glu Pro
                350
                                     355
                                                          360
Glu Ile His Gln Thr Pro Asp Trp Glu Ile Val Phe Glu Asp Asn
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                                     370
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Pro Gly Arg Leu Asn Glu Arg Arg Phe Gly Tyr
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Met Cys Ser Arg Lys Lys Ala Glu Phe Ile Lys Gly Ser His Lys
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Cys Asn Val Cys Ser Arg Thr Phe Phe Ser Glu Asn Gly Leu Arg
                 20
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Glu His Leu Gln Thr His Arg Gly Pro Ala Lys His Tyr Met
                                                         Cys
                                                           45
                 35
                                      40
Pro Ile Cys Gly Glu Arg Phe Pro Ser Leu Leu Thr Leu Thr
                                                         Glu
                 50
                                      55
                                                           60
His Lys Val Thr His Ser Lys Ser Leu Asp Thr Gly Thr Cys
                 65
Ile Cys Lys Met Pro Leu Gln Ser Glu Glu Glu Phe Ile Glu His
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Cys Gln Met His Pro Asp Leu Arg Asn Ser Leu Thr Gly Phe Arg
                 95
                                     100
                                                          105
Cys Val Val Cys Met Gln Thr Val Thr Ser Thr Leu Glu Leu Lys
                110
                                     115
                                                          120
Ile His Gly Thr Phe His Met Gln Lys Leu Ala Gly Ser Ser Ala
                                     130
                125
                                                          135
Ala Ser Ser Pro Asn Gly Gln Gly Leu Gln Lys Leu Tyr Lys Cys
                                                          150
                140
                                     145
Ala Leu Cys Leu Lys Glu Phe Arg Ser Lys Gln Asp Leu Val Lys
                                     160
                                                          165
                155
Leu Asp Val Asn Gly Leu Pro Tyr Gly Leu Cys Ala Gly Cys Met
                                     175
                170
Ala Arg Arg Ala Asn Gly Gln Val Gly Gly Leu Ala Pro Pro Glu
                185
                                     190
                                                          195
Pro Ala Asp Arg Pro Cys Ala Gly Leu Arg Cys Pro Glu Cys Ser
                200
                                     205
                                                          210
Val Lys Phe Glu Ser Ala Glu Asp Leu Glu Ser His Met Gln Val
                215
                                     220
                                                          225
Asp His Arg Asp Leu Thr Pro Glu Thr Ser Gly Pro Arg Lys Gly
                                     235
                                                          240
                230
Thr Gln Thr Ser Pro Val Pro Arg Lys Lys Thr Tyr Gln Cys Ile
                                                          255
                245
                                     250
Lys Cys Gln Met Thr Phe Glu Asn Glu Arg Glu Ile Gln Ile His
                                     265
                260
Asp Ala Asn His Met Ile Glu Glu Gly Ile Asn His Glu Cys Lys
                275
                                     280
                                                          285
Leu Cys Asn Gln Met Phe Asp Ser Pro Ala Lys Leu Cys His
                290
                                     295
                                                          300
Leu Ile Glu His Ser Phe Glu Gly Met Gly Gly Thr Phe Lys Cys
                305
                                     310
                                                          315
Pro Val Cys Phe Thr Val Phe Val Gln Ala Asn Lys Leu Gln Gln
                320
                                     325
                                                          330
His Ile Phe Ala Val His Gly Gln Glu Asp Lys Ile Tyr Asp Cys
                335
                                     340
Ser Gln Cys Pro Gln Lys Phe Phe Phe Gln Thr Glu Leu Gln Asn
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His Thr Met Ser Gln His Ala Gln
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Val Thr Phe Gln Asp Val Ala Ile Asp Phe Ser Lys Glu Glu Trp
Gly Phe Leu Asn Pro Ala Gln Arg Asp Leu Tyr Thr Thr Val Met
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                                      25
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Leu Glu Asn Tyr Gln Asn Leu Val Trp Leu Gly Leu Ser Ile Ser
                  35
                                      40
                                                           45
Lys Ser Val Ile Ser Leu Leu Glu Lys Arg Lys Leu Pro Trp Ile
                                      55
                                                           60
                  50
Met Ala Lys Glu Glu Ile Arg Gly Pro Leu Pro Asp Val Pro Gly
                                      70
                  65
Ala Glu Ile Lys Glu Leu Ser Ala Lys Arg Ala Ile Asn Glu Val
                                      85
                                                           90
                  80
Leu Ser Gln Phe Asp Thr Val Ile Lys Cys Thr Arg Asn Val Cys
                                                          105
                                     100
Lys Glu Cys Gly Asn Leu Tyr Cys His Asn Met Gln Leu Thr Leu
                 110
                                     115
His Lys Arg Asn His Thr Gln Lys Lys Cys Asn Gln Cys Leu Asp
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130
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                125
Cys Gly Lys Tyr Phe Thr Arg Gln Ser Pro Leu Ile Gln His Gln
                140
                                     145
                                                          150
Arg Ile His Thr Gly Glu Arg Pro Tyr Lys Cys Asn Glu Cys Ile
                                     160
                                                          165
                155
Lys Thr Phe Asn Gln Arg Ala His Leu Thr
                170
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Thr Gln Glu Val Glu Val Ala Val Ser Leu Asp Cys Cys His Cys
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Thr Pro Ala Trp Val Thr Ala Glu Leu Cys Leu Lys Lys Met
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                                                           30
Trp Asn Ser Phe Leu Gln Met Phe Ser Asn Ser Ile Pro Ser Ser
                                      40
                 35
Val Cys Arg Tyr Met Tyr Ala Ile Ile Leu Gln Val Ile His Val
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Asp Cys Ile Gly Asn Tyr Arg Lys Asp Tyr Ile Gly Leu Phe Arg
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Lys Tyr Phe
<210> 311
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Glu Gln Ser Ser Val Gln Gly Arg Ser Val Glu Val Leu Thr Val
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Gln Val Gln Met Leu Arg Asn Met Ser Pro Ala Met Ser Phe Leu
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Met Leu Gln Pro Cys Val Asp Gln Ser Ala Ser Gly Cys Asp Trp
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Ser Arg Ala Cys Arg Leu Leu Gln Ser Tyr Phe Ala Gly Val Gly
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Glu
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Gly Leu Phe Gln Cys Ile His Gln Val Thr Glu Val Gly Gln Lys
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Val Ala Thr Val Leu Leu Phe Tyr Gly Tyr Tyr Lys Cys Thr Gly 20 25 30
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Thr Leu Lys Ile Thr Cys Leu Tyr Asn Val Ile Leu Tyr Lys Val
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35
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Cys Ser Pro Gly Ser Asp Gln Pro Asp Val Cys Tyr Asp Pro Ser
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Glu Leu Pro Met Thr Thr Val Phe Lys Ile Arg Leu Arg
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Cys Cys Ser Phe Lys Phe His Phe Asp Leu Ser Trp Glu Ile Leu
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Trp Pro Ile Ile Pro Trp Met Leu Lys Met Val Leu Thr Glu Asn
                                                          30
                                      25
                 20
Pro Asn Gln Glu Ile Ala Thr Ser Leu Glu Phe Leu Leu Gln
                 35
                                      40
                                                          45
Asn Ser Pro Gly Ser Leu Arg Ala Gln Gln Arg Met Ser Tyr
                                                         Tyr
                 50
                                      55
                                                           60
Gly Ser Ser Tyr His Ile Ile Asn Ala Asp Ala Lys Tyr Pro Gly
                                      70
Tyr Pro Pro Glu His Ile Ile Ala Glu Lys Arg Arg Ala Arg Arg
                 80
                                      85
                                                          9.0
Arg Leu Leu His Lys Asp Gly Ser Cys Asn Val Tyr Phe Lys His
                                                          105
                 95
                                     100
Ile Phe Gly Glu Trp Gly Ser Tyr Val Val Asp Ile Phe Thr Thr
                                                          120
                                     115
                110
Leu Val Asp Thr Lys Trp Arg His Met Phe Val Ile Phe Ser Leu
                                     130
                125
Ser Tyr Ile Leu Ser Trp Leu Ile Phe Gly Ser Val Phe Trp Leu
                140
                                     145
                                                          150
Ile Ala Phe His His Gly Asp Leu Leu Asn Asp Pro Asp Ile Thr
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                                     160
                                                          165
Pro Cys Val Asp Asn Val His Ser Phe Thr Gly Ala Phe Leu Phe
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                170
Ser Leu Glu Thr
<210> 314
<211> 219
<212> PRT
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Leu Ser Gly His Val Gln Thr Leu Glu Ser Pro Pro Gln Cys Ser
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Pro Ala Pro Gly Gln Pro Asn Phe Cys Leu Leu Asp Gly Asp Gln
                                                           30
                                      25
                 20
Val Ala Ala Gly Ala Gly Ala Val Pro Ala Gly Val Glu Cys
                                      40
Leu Gly Leu Leu Val Arg Gln Arg Gly Arg Gly Gln Lys Cys Leu
                 50
Pro Ser Leu Pro Gln Thr Gln Glu Ala Gly Pro Ala Ala Ala Leu
                 65
                                      70
Arg Pro Arg Ser Thr Pro Cys Phe Val Tyr Gln Pro Ala Ile Arg
                 80
                                      85
                                                           90
Glu Ala Asn Gly Ile Val Glu Cys Gly Pro Cys Gln Lys Val Phe
                 95
                                     100
                                                          105
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Val Val Gln Gln Ile Pro Asn Ser Asn Leu Leu Leu Val Thr
                                                         120
                110
                                     115
Asp Pro Thr Cys Asp Cys Ser Ile Phe Pro Pro Val Leu Gln Glu
                                                         135
                125
                                     130
Ala Thr Glu Val Lys Tyr Asn Ala Ser Val Lys Cys Asp Arg Met
                140
                                     145
                                        Ser Cys His Ala Phe
Arg Ser Gln Lys Leu Arg Arg Arg Pro Asp
                                     160
                155
His Pro Glu Val Arg Val Glu Ala Asp Arg Gly Trp Ala Gly Phe
                170
                                     175
                                                          180
Ser Ser Pro Asn Pro Leu Cys Leu Gly Leu Cys Pro Cys Arg Gln
                185
                                     190
                                                          195
Glu His Ile Gly Met Pro Met Asn Thr Pro Val Pro Val Leu Leu
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                                     205
                                                          210
Gly Gly Asn Ile Arg Val Tyr Ala Leu
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<210> 315
<211> 1603
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Ala Ala Glu Pro Ala Pro Ser Ala Pro Ser Pro Ala Pro Ser Arg
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Val Arg Ala Lys Gln Gln Pro Pro Thr Pro Gly Pro Gly Arg
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Thr Ser Ser Phe Pro Thr Gly Asn Val Arg Arg Ala Cys Ala Gln
                                      40
                                                           45
                 35
Lys Gln Asp Glu Lys Met Ala Asn Phe Leu Leu Pro Arg Gly Thr
                                      55
                  50
Ser Ser Phe Arg Arg Phe Thr Arg Glu Ser Leu Ala Ala Ile Glu
                                      70
                                                           75
                  65
Lys Arg Met Ala Glu Lys Gln Ala Arg Gly Ser Thr Thr Leu Gln
                                      85
                                                           90
Glu Ser Arq Glu Gly Leu Pro Glu Glu Glu Ala Pro Arg Pro Gln
                  95
                                     100
                                                          105
Leu Asp Leu Gln Ala Ser Lys Lys Leu Pro Asp Leu Tyr Gly Asn
                                                          120
                 110
                                     115
Pro Pro Gln Glu Leu Ile Gly Glu Pro Leu Glu Asp Leu Asp Pro
                                     130
                 125
Phe Tyr Ser Thr Gln Lys Thr Phe Ile Val Leu Asn Lys Gly Lys
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                                     145
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Thr Ile Phe Arg Phe Ser Ala Thr Asn Ala Leu Tyr Val Leu Ser
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                                     160
                                                          165
Pro Phe His Pro Val Arg Arg Ala Ala Val Lys Ile Leu Val His
                 170
                                     175
                                                          180
Ser Leu Phe Asn Met Leu Ile Met Cys Thr Ile Leu Thr Asn Cys
                                                          195
                 185
                                     190
Val Phe Met Ala Gln His Asp Pro Pro Pro Trp Thr Lys Tyr Val
                                     205
                                                          210
                 200
Glu Tyr Thr Phe Thr Ala Ile Tyr Thr Phe Glu Ser Leu Val Lys
                 215
                                     220
                                                          225
Ile Leu Ala Arg Ala Phe Cys Leu His Ala Phe Thr Phe Leu Arg
                                                          240
                 230
                                     235
Asp Pro Trp Asn Trp Leu Asp Phe Ser Val Ile Ile Met Ala Tyr
                 245
                                     250
                                                          255
Thr Thr Glu Phe Val Asp Leu Gly Asn Val Ser Ala Leu Arg Thr
                                                          270
                 260
                                     265
Phe Arg Val Leu Arg Ala Leu Lys Thr Ile Ser Val Ile Ser
                                                          Gly
                                                          285
                 275
                                     280
Leu Lys Thr Ile Val Gly Ala Leu Ile Gln Ser Val Lys Lys Leu
```

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Ala Asp Val Met Val Leu Thr Val Phe Cys Leu Ser Val Phe Ala
                 305
                                     310
Leu Ile Gly Leu Gln Leu Phe Met Gly Asn Leu Arg His Lys Cys
                 320
                                     325
                                                          330
Val Arg Asn Phe Thr Ala Leu Asn Gly Thr Asn Gly Ser Val Glu
                                      340
                                                           345
                 335
Ala Asp Gly Leu Val Trp Glu Ser Leu Asp Leu Tyr Leu Ser Asp
                 350
                                     355
                                                          360
Pro Glu Asn Tyr Leu Leu Lys Asn Gly Thr Ser Asp Val Leu Leu
                                     370
                                                           375
                 365
Cys Gly Asn Ser Ser Asp Ala Gly Thr Cys Pro Glu Gly Tyr Arg
                                     385
                 380
Cys Leu Lys Ala Gly Glu Asn Pro Asp His Gly Tyr Thr Ser Phe
                 395
                                      400
Asp Ser Phe Ala Trp Ala Phe Leu Ala Leu Phe Arg Leu Met Thr
                 410
                                      415
                                                           420
Gln Asp Cys Trp Glu Arg Leu Tyr Gln Gln Thr Leu Arg Ser Ala
                 425
                                      430
                                                           435
Gly Lys Ile Tyr Met Ile Phe Phe Met Leu Val Ile Phe Leu Gly
                                      445
                 440
Ser Phe Tyr Leu Val Asn Leu Ile Leu Ala Val Val Ala Met Ala
                 455
                                      460
                                                           465
Tyr Glu Glu Gln Asn Gln Ala Thr Ile Ala Glu Thr Glu Glu Lys
                                                           480
                 470
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Glu Lys Arg Phe Gln Glu Ala Met Glu Met Leu Lys Lys Glu His
                                                           495
                                      490
                 485
Glu Ala Leu Thr Ile Arg Gly Val Asp Thr Val Ser Arg Ser Ser
                 500
                                      505
Leu Glu Met Ser Pro Leu Ala Pro Val Asn Ser His Glu Arg Arg
                 515
                                      520
                                                           525
Ser Lys Arg Arg Lys Arg Met Ser Ser Gly Thr Glu Glu Cys Gly
                 530
                                      535
                                                           540
Glu Asp Arg Leu Pro Lys Ser Asp Ser Glu Asp Gly Pro Arg Ala
                                                           555
                                      550
                 545
Met Asn His Leu Ser Leu Thr Arg Gly Leu Ser Arg Thr Ser Met
                                      565
                                                           570
                 560
Lys Pro Arg Ser Ser Arg Gly Ser Ile Phe Thr Phe Arg Arg Arg
                                                           585
                 575
                                      580
Asp Leu Gly Ser Glu Ala Asp Phe Ala Asp Asp Glu Asn Ser Thr
                 590
                                      595
                                                           600
Ala Arg Glu Ser Glu Ser His His Thr Ser Leu Leu Val Pro Trp
                 605
                                      610
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Pro Leu Arg Arg Thr Ser Ala Gln Gly Gln Pro Ser Pro Gly Thr
                                                           630
                 620
                                      625
 Ser Ala Pro Gly His Ala Leu His Gly Lys Lys Asn Ser Thr Val
                                      640
                 635
Asp Cys Asn Gly Val Val Ser Leu Leu Gly Ala Gly Asp Pro Glu
                 650
                                      655
Ala Thr Ser Pro Gly Ser His Leu Leu Arg Pro Val Met Leu Glu
                 665
                                      670
                                                           675
His Pro Pro Asp Thr Thr Pro Ser Glu Glu Pro Gly Gly Pro
                 680
                                      685
                                                           690
Gln Met Leu Thr Ser Gln Ala Pro Cys Val Asp Gly Phe Glu Glu
                                                           705
                                      700
                 695
Pro Gly Ala Arg Gln Arg Ala Leu Ser Ala Val Ser Val Leu Thr
                                      715
                                                           720
                 710
 Ser Ala Leu Glu Glu Leu Glu Glu Ser Arg His Lys Cys Pro Pro
                                                           735
                                      730
                 725
Cys Trp Asn Arg Leu Ala Gln Arg Tyr Leu Ile Trp Glu Cys
                                                          Cys
                                      745
                 740
 Pro Leu Trp Met Ser Ile Lys Cln Cly Val Lys Leu Val Val Met
                 755
                                      760
                                                           765
Asp Pro Phe Thr Asp Leu Thr Ile Thr Met Cys Ile Val Leu Asn
                 770
                                      775
                                                           780
 Thr Leu Phe Met Ala Leu Glu His Tyr Asn Met Thr Ser Glu Phe
                                                           795
                 785
                                      790
 Glu Glu Met Leu Gln Val Gly Asn Leu Val Phe Thr Gly Ile Phe
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800
Thr Ala Glu Met Thr Phe Lys Ile Ile Ala Leu Asp Pro Tyr Tyr
                815
                                     820
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Tyr Phe Gln Gln Gly Trp Asn Ile Phe Asp Ser Ile Ile Val Ile
                830
                                     835
                                                          840
Leu Ser Leu Met Glu Leu Gly Leu Ser Arg Met Ser Asn Leu Ser
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                845
                                     850
Val Leu Arg Ser Phe Arg Leu Leu Arg Val
                                        Phe Lys Leu Ala Lys
                                                          870
                860
                                     865
   Trp Pro Thr Leu Asn Thr Leu Ile Lys
                                         Ile Ile Gly Asn Ser
                                                          885
                875
                                     880
Val Gly Ala Leu Gly Asn Leu Thr Leu Val Leu Ala Ile Ile Val
                890
                                     895
                                                          900
Phe Ile Phe Ala Leu Val Gly Lys Gln Leu Leu Gly Glu Asn Tyr
                                                          915
                905
                                     910
Arg Asn Asn Arg Lys Asn Ile Ser Ala Pro His Glu Asp Trp Pro
                920
                                     925
                                                          930
Arg Trp His Met His Asp Phe Phe His Ser Phe Leu Ile Val Phe
                935
                                     940
                                                          945
Arg Ile Leu Cys Gly Glu Trp Ile Glu Asn Met Trp Ala Cys Met
                950
                                     955
                                                          960
Glu Val Gly Gln Lys Ser Ile Cys Leu Ile Leu Phe Leu Thr Val
                                                          975
                965
                                     970
Met Val Leu Gly Asn Leu Val Val Leu Asn Leu Phe Ile Ala Leu
                                     985
                                                          990
                980
Leu Leu Asn Ser Phe Ser Ala Asp Asn Leu Thr Ala Pro Glu Asp
                995
                                    1000
                                                         1005
Asp Gly Glu Val Asn Asn Leu Gln Val Ala Leu Ala Arg Ile Gln
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Arg Gly Leu Arg Phe Val Lys Arg Thr Thr
                                         Trp Asp Phe Cys Cys
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Gly Leu Leu Arg His Arg Pro Gln Lys Pro Ala Ala Leu Ala Ala
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Gln Gly Gln Leu Pro Ser Cys Ile Ala Thr Pro Tyr Ser Pro Pro
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Pro Pro Glu Thr Glu Lys Val Pro Pro Thr Arg Lys Glu Thr Gln
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                                    1075
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Phe Glu Glu Gly Glu Gln Pro Gly Gln Gly
                                         Thr Pro Gly Asp Pro
               1085
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                                                         1095
Glu Pro Val Cys Val Pro Ile Ala Val Ala Glu Ser Asp Thr Asp
               1100
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Asp Gln Glu Glu Asp Glu Glu Asn Ser Leu Gly Thr Glu Glu Glu
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                                    1120
                                                         1125
Ser Ser Lys Gln Gln Glu Ser Gln Pro Val Ser Gly Trp Pro Arg
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                                                         1140
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Gly Pro Pro Asp Ser Arg Thr Trp Ser Gln Val Ser Ala Thr Ala
               1145
                                    1150
Ser Ser Glu Ala Glu Ala Ser Ala Ser Gln Ala Asp Trp Arg Gln
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               1160
                                    1165
Gln Trp Lys Ala Glu Pro Gln Ala Pro Gly
                                         Cys Gly Glu Thr Pro
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                                    1180
                                                         1185
Glu Asp Ser Cys Ser Glu Gly Ser Thr Ala Asp Met Thr Asn Thr
               1190
                                    1195
                                                         1200
Ala Glu Leu Leu Glu Gln Ile Pro Asp Leu Gly Gln Asp Val Lys
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               1205
                                    1210
                    Phe Thr Glu Gly Cys
                                         Val Arg Arg Cys Pro
Asp Pro Glu Asp Cys
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               1220
Cys Cys Ala Val Asp
                    Thr Thr Gln Ala Pro Gly Lys Val Trp Trp
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                    Cys Tyr His Ile Val Glu His Ser Trp Phe
Arg Leu Arg Lys Thr
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                                    1255
                                                         1260
Glu Thr Phe Ile Ile Phe Met Ile Leu Leu Ser Ser Gly Ala Leu
               1265
                                    1270
                                                         1275
Ala Phe Glu Asp Ile Tyr Leu Glu Glu Arg Lys Thr Ile Lys Val
               1280
                                    1285
                                                         1290
Leu Leu Glu Tyr Ala Asp Lys Met Phe Thr Tyr Val Phe Val Leu
               1295
                                    1300
                                                         1305
```

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Glu Met Leu Leu Lys Trp Val Ala Tyr Gly Phe Lys Lys Tyr Phe
               1310
                                   1315
Thr Asn Ala Trp Cys Trp Leu Asp Phe Leu Ile Val Asp Val Ser
               1325
                                                        1335
                                   1330
Leu Val Ser Leu Val Ala Asn Thr Leu Gly Phe Ala Glu Met Gly
               1340
                                   1345
                                                        1350
Pro Ile Lys Ser Leu Arg Thr Leu Arg Ala Leu Arg Pro Leu Arg
                                                        1365
               1355
                                   1360
Ala Leu Ser Arg Phe Glu Gly Met Arg Val Val Asn Ala Leu
                                   1375
                                                        1380
               1370
Val Gly Ala Ile Pro Ser Ile Met Asn Val Leu Leu Val Cys Leu
                                   1390
                                                        1395
               1385
Ile Phe Trp Leu Ile Phe Ser Ile Met Gly Val Asn Leu Phe Ala
               1400
                                   1405
                                                        1410
Gly Lys Phe Gly Arg Cys Ile Asn Tyr Thr Asp Gly Glu Phe Ser
                                                        1425
               1415
                                    1420
Leu Val Pro Leu Ser Ile Val Asn Asn Lys Ser Asp Cys Lys Ile
                                    1435
               1430
Gln Asn Ser Thr Gly Ser Phe Phe Trp Val Asn Val Lys Val Asn
               1445
                                   1450
                                                        1455
Phe Asp Asn Val Ala Met Gly Tyr Leu Ala Leu Leu Gln Val Ala
                                                        1470
               1460
                                    1465
Thr Phe Lys Gly Trp Met Asp Ile Met Tyr Ala Ala Val Asp Ser
                                                        1485
                                    1480
               1475
Arg Glu Val Asn Met Gln Pro Lys Trp Glu Asp Asn Val Tyr Met
                                   1495
               1490
Tyr Leu Tyr Phe Val Ile Phe Ile Phe Gly Gly Phe Phe Thr
                                    1510
               1505
                                                        1515
Leu Asn Leu Phe Val Gly Val Ile Ile Asp Asn Phe Asn Gln Gln
               1520
                                    1525
                                                        1530
Lys Lys Leu Gly Gly Gln Asp Ile Phe Met Thr Glu Glu Gln
               1535
                                    1540
                                                        1545
Lys Lys Tyr Tyr Asn Ala Met Lys Lys Leu Gly Ser Lys Lys Pro
                                                        1560
                                    1555
               1550
Gln Lys Pro Ile Pro Arg Pro Leu Asn Lys
                                        Phe Gln Gly Phe Val
                                   1570
                                                        1575
               1565
Phe Asp Ile Val Thr Arg Gln Ala Phe Asp Ile Thr Ile Met Val
                                                        1590
               1580
                                    1585
Leu Ile Cys Leu Asn Met Val His His Asp Gly Gly Asp
               1595
                                    1600
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Gly Gly Ala Gly Val Arg Gly Ala Gly Trp Arg Leu Gln Gln Arg
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                                      10
Gly Arg Gly Ala Arg Ala Pro Pro Ala Arg Gln Ala Arg Gln Gly
                 20
                                                           30
Gly Pro Arg Ala Arg Pro Leu Glu Gln Gln Gly Val Arg
                                                         Ala
                                      40
                                                           45
Glu Arg Gly Arg Gly Asp His Trp Ala Trp Ala Thr Cys Gly Ala
                                      55
                                                           60
Ser Pro Thr Leu Cys Tyr Lys Asn Gly Gly Gly Ala Phe Leu Ile
                                                           75
                 65
                                      70
Pro Tyr Val Val Phe Phe Ile Cys Cys Gly Ile Pro Val Phe Phe
                                                           90
                                      85
                 80
Leu Glu Thr Ala Leu Gly Gln Phe Thr Ser Glu Gly Gly Ile Thr
                                     100
                                                          105
                 95
Cys Trp Arg Lys Val Cys Pro Leu Phe Glu Gly Ile Gly Tyr Ala
                110
```

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Thr Gln Val Ile Glu Ala His Leu Asn Val Tyr Tyr Ile Ile Ile
                125
                                     130
Leu Ala Trp Ala Ile Phe Tyr Leu Ser Asn Cys Phe Thr Thr Glu
                140
                                     145
                                                          150
Leu Pro Trp Ala Thr Cys Gly His Glu Trp Asn Thr Glu Asn Cys
                155
                                     160
                                                          165
Val Glu Phe Gln Lys Leu Asn Val Ser Asn Tyr Ser His Val Ser
                                     175
                                                          180
                170
Leu Gln Lys Cys His Leu Pro Cys His Gly Val Leu Gly Ala Pro
                185
Gly Pro Gly His Leu
                200
<210> 317
<211> 329
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Arg Thr Gly Gly Gly Gly Ser Gln Ser Pro Ala Pro Asp Gly
                                      10
                                                           15
Glu Arg Thr Met His Cys Leu Gly Ala Glu Tyr Leu Val Ser Ala
                                                           3.0
                                      25
Asp Gly Ala Pro Arg Gln Arg Glu Trp Arg Pro Gln Ile Tyr Arg
                                      40
                 35
Lys Cys Thr Asp Thr Ala Trp Leu Phe Leu Phe Phe Leu Phe Trp
                                      55
                                                           60
                 50
Thr Gly Leu Val Phe Ile Met Gly Tyr Ser Val Val Ala Gly Ala
                                      70
Ala Gly Arg Leu Leu Phe Gly Tyr Asp Ser Phe Gly Asn Met Cys
                                                           90
                 80
                                      85
Gly Lys Lys Asn Ser Pro Val Glu Gly Ala Pro Leu Ser Gly Gln
                                                          105
                 95
                                     100
Asp Met Thr Leu Lys Lys His Val Phe Phe Met Asn Ser Cys Asn
                                     115
                110
Leu Glu Val Lys Gly Thr Gln Leu Asn Arg Met Ala Leu Cys Val
                                     130
                                                          135
                125
Ser Asn Cys Pro Glu Glu Gln Leu Asp Ser Leu Glu Glu Val Gln
                140
                                     145
                                                          150
Phe Phe Ala Asn Thr Ser Gly Ser Phe Leu Cys Gly Tyr Ser Leu
                155
                                     160
                                                          165
Asn Ser Phe Asn Tyr Thr His Ser Pro Lys Ala Asp Ser Leu Cys
                                                          180
                                     175
                170
Pro Arg Leu Pro Val Pro Pro Ser Lys Ser Phe Pro Leu Phe Asn
                185
                                     190
                                                          195
Arg Cys Val Pro Gln Thr Pro Glu Cys Tyr Ser Leu Phe Ala Ser
                                     205
                                                          210
                200
Val Leu Ile Asn Asp Val Asp Thr Leu His Arg Ile Leu Ser Gly
                215
                                     220
                                                          225
Ile Met Ser Gly Arg Asp Thr Ile Leu Gly Leu Cys Ile Leu Ala
                230
                                     235
                                                          240
Leu Ala Leu Ser Leu Ala Met Met Leu Thr Val Gln Ile His Thr
                                                          255
                                     250
                245
Pro Pro Phe Trp Phe Thr Phe Ser Phe His Trp Leu Phe Trp Asp
                                     265
                                                          270
                260
Cys Cys Leu Val Cys Gly Val Leu Trp Trp Leu Tyr Tyr Asp
                                                          Tyr
                275
                                     280
                                                          285
Thr Asn Asp Leu Ser Ile Glu Leu Asp Thr Glu Gln Gly Lys
                                                          Tyr
                290
                                     295
                                                          300
Glu Val Arg Ala Gly Val Cys Tyr Arg Asn Pro Gln Gly Ile
                                                          Thr
                305
                                     310
Ala Asp Ala Ala Arg Leu Asp Ile Leu Phe Ser Glu Arg Glu
                320
                                     325
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<210> 318
<211> 256
<212> PRT
<213> Homo sapiens
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<400> 318
His Leu Ala Ala Ala Met Ser Arg Arg Tyr Asp Ser Arg Thr
                                      10
Thr Ile Phe Ser Pro Glu Gly Arg Leu Tyr Gln Val Glu Tyr Ala
                                                           30
                 20
Met Glu Ala Ile Gly Asn Ala Gly Ser Ala Leu Gly Ile Leu Ala
                                      40
Ala Asp Gly Val Val Leu Val Gly Glu Lys Lys Val Thr Ser Lys
                 50
                                      55
                                                           60
Leu Leu Gln Thr Ser Arg Ser Ala Glu Lys Met Tyr Lys Ile Asp
                 65
                                      70
                                                           75
Ser His Leu Ala Cys Ala Val Ala Gly Ile Met Ser Asp Ala Asn
                 80
                                      85
                                                           90
Ile Leu Ile Asn Thr Ala Arg Leu His Ala Gln Arg Tyr Ala Leu
                                     100
                                                          105
                 95
Ser Tyr Gln Glu Pro Ile Pro Val Glu Gln Leu Val Gln Ser Leu
                                     115
                                                          120
                110
Cys Asp Thr Lys Gln Gly Tyr Thr Gln Phe Gly Gly Leu Arg Pro
                125
                                     130
                                                          135
Phe Gly Val Ser Phe Leu Phe Ala Gly Trp Asp Lys His His Gly
                140
                                     145
                                                          150
Phe Gln Leu Tyr Met Ser Asp Pro Ser Gly Asn Tyr Gly Gly Trp
                155
                                     160
                                                          165
Lys Ala Ala Ala Val Gly Ala Asn Ser Gln Ala Ala Gln Ser Met
                170
                                     175
                                                          180
Leu Lys Gln Asp Tyr Lys Asp Ala Leu Thr Arg Glu Glu Ala Val
                185
                                     190
                                                          195
Gly Leu Ala Leu Lys Val Leu Ser Lys Thr Met Asp Ser Thr Ser
                200
                                     205
                                                          210
Leu Thr Ala Glu Lys Leu Glu Leu Ala Glu Val Phe Leu Gln Pro
                215
                                     220
                                                          225
Asp Thr Gly Glu Val Gln Tyr Gln Val Cys Ser Pro Glu Ala Leu
                230
                                     235
                                                          240
Gly Lys Leu Leu Ala Asn Ser Gly Leu Thr Gln Pro Thr Pro Glu
                245
Ala
<210> 319
<211> 76
<212> PRT
<213> Homo sapiens
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<400> 319
Ala Val Ser Phe Arg Arg Leu Leu Gln Thr Trp Ser Thr Pro Pro
                                      10
                                                           15
Cys Ser Ser Thr Ser Arg Leu Met Ala Ser Leu Trp Val Ala Val
                 20
                                      25
                    Leu Ala Asp Lys Val Pro Lys Thr Ala Glu
Trp Leu Arg Lys Trp
                                      40
                 35
Asn Phe Arg Ala Leu Ser Thr Gly Glu Lys Gly Phe Gly Tyr Asn
                 50
                                      55
Gly Phe Leu Leu Ser Gln Asn Tyr Ser Arg Ile His Val Pro Gly
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Trp

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<210> 320
<211> 276
<212> PRT
<213> Homo sapiens
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Thr Asn Ile Lys Ile Thr Met Lys Val Leu Gly Val Thr Lys Asp
                                      10
                                                           15
Ser Gly Asp Glu Asp Leu Lys Lys Ala Tyr Arg Lys Leu Ala Leu
                                                           30
                                      25
                 20
Lys Phe His Pro Asp Lys Asn His Ala Pro Gly Ala Thr Asp
                                      40
                 35
Phe Lys Lys Ile Gly Asn Ala Tyr Ala Val Leu Ser Asn Pro Glu
                 50
                                      55
                                                           60
Lys Arg Lys Gln Tyr Asp Leu Thr Gly Asn Glu Glu Gln Ala Cys
                                      70
Asn His Gln Asn Asn Gly Arg Phe Asn Phe His Arg Gly Cys Glu
                 80
                                      85
                                                           90
Ala Asp Ile Thr Pro Glu Asp Leu Phe Asn Ile Phe Phe Gly Gly
                 95
                                     100
                                                          105
Gly Phe Pro Ser Gly Ser Val His Ser Phe Ser Asn Gly Arg Ala
                110
                                     115
                                                          120
Gly Tyr Ser Gln Gln His Gln His Arg His Ser Gly His Glu Arg
                125
                                     130
                                                          135
Glu Glu Glu Arg Gly Asp Gly Gly Phe Ser Val Phe Ile Gln Leu
                140
                                     145
Met Pro Ile Ile Val Leu Ile Leu Val Ser Leu Leu Ser Gln Leu
                155
                                     160
                                                          165
                                         Pro Arg Ser Gly
Met Val Ser Asn Pro Pro Tyr Ser Leu Tyr
                                                          Thr
                170
                                     175
                                                          180
Gly Gln Thr Ile Lys Met Gln Thr Glu Asn Leu Gly Val Val
                                                          195
                                     190
                185
Tyr Val Asn Lys Asp Phe Lys Asn Glu Tyr
                                         Lys Gly Met Leu Leu
                200
                                     205
                                         Tyr Val Thr Asn Ile
Gln Lys Val Glu Lys Ser Val Glu Glu Asp
                215
                                     220
                                                          225
Arg Asn Asn Cys Trp Lys Glu Arg Gln Gln Lys Thr Asp Met Gln
                230
                                     235
                                                          240
Tyr Ala Ala Lys Val Tyr Arg Asp Asp Arg Leu Arg Arg Lys Ala
                                     250
                245
                                                          255
Asp Ala Leu Ser Met Asp Asn Cys Lys Glu Leu Glu Arg Leu Thr
                260
                                     265
Ser Leu Tyr Lys Gly Gly
                275
<210> 321
<211> 115
<212> PRT
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<400> 321
Gln Ala Leu Leu Gln Ser His Pro Glu Ala Asp Trp Ser Thr His
                                      10
                                                           15
Ser Arg Ser Met Arg Lys Leu Ile Val Arg Phe Ile Phe Leu Lys
                                      25
                                                           30
Phe Trp Thr Tyr Thr Val Arg Ala Ser Thr Asp Leu Thr Gln Thr
```

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35
Gly Asp Cys Ser Gln Cys Ile His Gln Val Thr Glu Val Gly Gln
                 50
                                      55
                                                           60
Gln Ile Lys Thr Ile Phe Leu Phe Tyr Ser Tyr Tyr Glu Cys Val
                                      70
                                                           75
                 65
Glu Thr Ile Lys Lys Leu Val Cys Ile Met Pro Leu Ser Thr
                                                         Ara
                                                           90
                 80
                                      85
Tyr Val Ala Arg Glu Met Thr Asp Leu Met Arg Val Ile Thr
                                                         His
                 95
                                     100
                                                          105
Leu Ser Pro Pro Gln Pro Pro Phe Leu Lys
                110
<210> 322
<211> 227
<212> PRT
<213> Homo sapiens
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<223> Incyte ID No: LI:373637.1.orf3:2000FEB01
<400> 322
His Pro Ala Trp Trp Thr Thr Thr Arg Cys Trp Thr Cys Pro Gly
                                                           15
                                         Thr Ala Ser Trp Arg
Arg Pro His Pro Arg Pro Ser Arg Arg Arg
                                      25
                 20
                                                           30
Ser Ser Gly Thr Pro Thr Lys Thr Leu Arg Thr Arg Arg Lys Arg
                 35
                                      40
                                                           45
Arg Gly Asp Ser Ser Arg Trp Pro Arg Pro Thr Arg Cys Cys Arg
                                                           60
                                      55
                 50
Thr Pro Arg Asn Ala Ile Ser Met Thr Ala Met Ala Arg Arg Gly
                                      70
                 65
Arg Arg Ala Ala Ala Gln Ala Ala Gly Pro Ser Arg Thr Pro Ser
                                      85
                                                           90
                 80
Ser Thr Ser Ser Ala Ser Ala Thr Gln Pro Thr Ser Ser Gly Ser
                 95
                                     100
                                                          105
Ser Ser Ala Ala Arg Thr His Ser Pro Leu Thr Ser Trp Glu Thr
                110
                                     115
                                                          120
Arg Trp Arg Ile Phe Trp Gly Gly Gln Arg Asn Cys Trp Gly Ser
                                     130
                                                          135
                 125
Arg Ser Arg Ala Ser Ala Pro Leu Phe Ser Ala Phe Ser Glu Phe
                140
                                     145
                                                          150
Pro Ala Phe Gly Gly Val Phe Ser Ser Phe Asp Thr Gly Phe Arg
                                     160
                                                          165
                 155
Ser Phe Gly Ser Leu Gly Ser Gly Leu Ser Ser Phe Cys Met
                                     175
                                                          180
                 170
Ser Tyr Gly Ser Asp Gly Thr Gly Ser Phe Lys Ser Met Ser Thr
                 185
                                     190
                                                          195
Ser Thr Glu Ile Val Asp Gly Lys Lys Ile Thr Thr Lys Arg Ile
                                     205
                                                          210
                 200
Ile Glu Asn Gly Gln Glu Arg Val Glu Val Glu Glu Asp Gly Glu
                                     220
                215
Leu Ser
<210> 323
<211> 100
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<223> Incyte ID No: LG:239368.2.orf3:2000MAY19
<400> 323
Glu Glu Ser Val Leu Arg Gly Lys Phe Leu Phe Thr Ser Gly Ile
                                       10
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Pro Arg Ala Ser Trp Val Asp Ser Gly Leu His Thr Gln Pro Gly
                 20
                                     25
Ser Pro Gly Ser Ala Ser Val Pro Pro Leu Ser Gly Pro Gly Cys
Gly Leu Gly Ala Arg Pro Ser Leu Ala Pro Gly Asn Ser Asp Val
                 50
                                      55
                                                          60
Phe Leu His Leu Leu Pro Leu Leu Arg Gly Pro Lys Pro Gly Lys
                                      70
                 65
Pro Ala Asp Ala Ser His Pro Glu Asn Cys Glu Gln Thr His
                                      85
                 80
Ala Ser Pro Thr Pro Glu Ser Ser Cys Cys
<210> 324
<211> 142
<212> PRT
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<400> 324
Asp Phe Trp Ala Lys Ile Tyr Leu Tyr Ala Leu Glu Gly Arg Lys
                                                           15
Tyr Arg Ser Ile Leu Gln Leu Val Lys Pro Trp Tyr Asp Glu Val
                                                           30
                 20
                                      25
Lys Asp Tyr Ala Phe Pro Tyr Pro Gln Asp Cys Asn Pro Arg Cys
                                      40
                 35
Pro Met Arg Cys Phe Gly Pro Met Cys Thr His Tyr Thr Gln Met
                                      55
                 50
Val Trp Ala Thr Ser Asn Arg Ile Gly Cys Ala Ile His Thr Cys
                 65
                                      70
                                                           75
Gln Asn Met Asn Val Trp Gly Ser Val Trp Arg Arg Ala Val Tyr
                 80
                                      85
Leu Val Cys Asn Tyr Ala Pro Lys Gly Asn Trp Ile Gly Glu Ala
                                                          105
                 95
                                     100
Pro Tyr Lys Val Gly Val Pro Cys Ser Ser Cys Pro Pro Ser Tyr
                                                          120
                110
                                     115
Gly Gly Ser Cys Thr Asp Asn Leu Cys Phe Pro Gly Val Thr Ser
                125
Asn Tyr Leu Tyr Trp Phe Lys
                140
<210> 325
<211> 263
<212> PRT
<213> Homo sapiens
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<223> Incyte ID No: LI:449393.1.orf3:2000MAY01
<400> 325
Ala Met Ser Leu Arg Val Leu Asn Pro Asn Ala Glu Val Leu Asn
                                      10
Lys Ser Ala Ala Leu His Met Asn Ile Asn Ala Ala Lys Gly Leu
                                      25
                 20
Gln Asp Val Leu Lys Thr Asn Leu Gly Pro Lys Gly Thr Ile Lys
                                                           45
                  35
                                      40
Met Leu Val Gly Gly Ala Gly Asp Leu Lys Leu Thr Lys Asp Gly
                                      55
                                                           60
Asn Thr Leu Leu Lys Glu Met Gln Ile Gln Asn Pro Thr Ala Ile
                  65
                                      70
                                                           75
Met Ile Ala Arg Thr Ala Val Ala Gln Asp Asp Thr Ser Gly Asp
                 80
                                      85
Gly Thr Thr Ser Thr Val Leu Phe Ile Gly Glu Leu Met Lys Gln
```

```
100
                                                          105
Ser Glu Arg Cys Ile Asp Glu Gly Thr His Pro Arg Phe Leu Val
                110
                                     115
                                                          120
Asp Gly Phe Asp Val Ala Lys Arg Ala Cys Leu Asp Phe Leu Asp
                                                          135
                                     130
                125
Lys Phe Lys Thr Pro Val Val Thr Gly Glu Glu Pro Asp Arg Asp
                                     145
                                                          150
                140
Thr Leu Lys Met Val Ala Arg Thr Thr Leu Arg Thr Lys Leu Tyr
                155
                                     160
                                                          165
Glu Gly Leu Ala Asp Gln Leu Thr Asp Ile Val Val Asn Ala Val
                                     175
                                                          180
                170
Leu Cys Ile Arg Lys Pro Asp Glu Pro Ile Asp Leu Phe Met Val
                                     190
                                                          195
                185
Glu Ile Met His Met Arg His Lys Phe Asp Val Asp Thr Arg Leu
                                     205
                                                          210
                200
Val Glu Gly Leu Val Leu Asp His Gly Ser Arg His Pro Asp Met
                215
                                     220
                                                          225
Lys Arg Arg Ala Glu Asn Cys Tyr Ile Leu Thr Cys Asn Val Ser
                                                          240
                230
                                     235
Leu Glu Tyr Glu Lys Ser Glu Ile Asn Ala Gly Phe Phe Tyr Ser
                                                          255
                245
Asn Ala Glu Gln Lys Lys Lys
                260
<210> 326
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Asp Glu Thr Arg Glu Asp Ala Met Ala Met Val Asp His Cys Leu
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                                         Val Lys Val Asp Leu
Lys Lys Ala Leu Trp Phe Gln Gly Arg Cys
                                      25
                 20
                    Lys Leu Val Leu Arg Ile Pro Asn Arg Gly
Ser Glu Lys Tyr Lys
                                                           45
                  35
                                      40
                                         Lys Arg Ser Tyr Ser
Ile Asp Leu Leu Lys
                    Lys Asp Lys Ser Arg
                                      55
                                                           60
Pro Asp Gly Lys Glu Ser Pro Ser Asp Lys Lys Ser Lys Thr
                                                         Asp
                                      70
                                                           75
Gly Ser Gln Lys Thr Glu Ser Ser Thr Glu Gly Lys Glu Gln Glu
                 80
                                      85
Glu Lys Ser Gly Glu Asp Gly Glu Lys Asp Thr Lys Asp Asp Gln
                                     100
                                                          105
                 95
Thr Glu Gln Glu Pro Asn Met Leu Leu Glu Ser Glu Asp Glu Leu
                                     115
                                                          120
                110
Leu Val Asp Glu Glu Glu Ala Ala Leu Leu Glu Ser Gly Ser
                                                          135
                 125
Ser Val Gly Asp Glu Thr Asp Leu Ala Asn Leu Gly Asp Val Ala
                                     145
                                                          150
                 140
Ser Asp Gly Lys Lys Glu Pro Ser Asp Lys Ala Val Lys Lys Asp
                 155
                                     160
                                                          165
Gly Ser Ala Ser Ala Ala Ala Lys Lys Lys Leu Lys Lys Arg Arg
                                                          180
                                     175
                 170
Phe Pro Gly Ser Met Glu Gly Phe Val Thr Leu Asp Glu Val Gly
                                                          195
                                     190
                 185
Asp Glu Glu Asp Ser Glu Leu Gln Lys Leu Arg Lys Ser Gly Met
                                                          210
                 200
                                     205
Ala Phe Lys Ser Gly Asp Lys Asn Asp Asp Gly Leu Val Glu Ile
                                     220
                                                          225
                 215
Lys Val Asp Lys Ile Glu Glu Leu Asp Gln Glu Asn Glu Ala Ala
                 230
                                     235
Leu Glu Asn Gly Ile Lys Asn Glu Glu Asn Thr Glu Pro Gly Ala
```

```
250
                245
Glu Ser Ser Glu Asn Ala Asp Asp Pro Asn Lys Asp Thr Ser Glu
                                                          270
                260
                                     265
Asn Ala Asp Gly Gln Ser Asp Glu Asn Lys Asp Asp Tyr Thr Ile
                275
                                     280
Pro Asp Glu Tyr Arg Ile Gly Pro Tyr Gln Pro Asn Val Pro Val
                                     295
                290
Gly Ile Asp Tyr Val Ile Pro Lys Thr Gly Phe Tyr Cys Lys Leu
                305
                                     310
                                                          315
Cys Ser Leu Phe Tyr Thr Asn Glu Glu Val Ala Lys Asn Thr His
                320
                                     325
                                                          330
Cys Ser Ser Leu Pro His Tyr Gln Lys Leu Lys Lys Phe Leu Asn
                335
                                     340
                                                          345
Lys Leu Ala Glu Glu Arg Arg Gln Lys Lys Glu Thr
                350
<210> 327
<211> 100
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<221> unsure
<222> 10, 18, 30
<223> unknown or other
<400> 327
Met Ile Ser Ser Asn Ser Pro Asn Leu Xaa Leu Trp Pro Ile Thr
                                      10
Thr Phe Xaa His Val Cys Thr Ser Cys Cys Ser Arg Leu Gln Xaa
                                      25
                                                           30
                 20
Pro Phe Ser Leu Ala Asp Phe Trp Lys Ser Asn Gly Arg Val Leu
                                                           45
                 35
                                      40
Gly Gly Arg Arg Leu Leu Tyr Ala Cys Glu Lys Glu Gln Ser Val
                 50
                                      55
Pro Thr Glu Gly Ser Ser Thr Thr Leu Leu Gln Asn Met Tyr Ile
                 65
                                      70
Ser Arg Leu Ser Ser His Leu Arg Phe Leu Cys Ser Cys Arg Leu
                 80
                                      85
Ile Asp Tyr Ser Ile Leu Leu Lys Arg Lys
                 95
<210> 328
<211> 303
<212> PRT
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<400> 328
Arg Glu Leu Lys Arg Phe Asn Ala Asp Asn Lys Leu Leu Thr
                                      10
Gly Thr Pro Leu Gln Asn Asn Leu Ser Glu Leu Trp Ser Leu Leu
                                                           30
                                      25
                 20
Asn Phe Leu Leu Pro Asp Val Phe Asp Asp Leu Lys Ser Phe Glu
                                                           45
                 35
                                      40
                                                          Ile
Ser Trp Phe Asp Ile Thr Ser Leu Ser Glu Thr Ala Glu Asp
                  50
                                      55
                                                           60
Ile Ala Lys Glu Arg Glu Gln Asn Val Leu His Met Leu His Gln
                                      70
Ile Leu Thr Pro Phe Leu Leu Arg Arg Leu Lys Ser Asp Val Ala
```

```
90
                 80
                                      85
Leu Glu Val Pro Pro Lys Arg Glu Val Val Val Tyr Ala Pro Leu
                 95
                                     100
Ser Lys Lys Gln Glu Ile Phe Tyr Thr Ala Ile Val Asn Arg Thr
                                     115
                110
Ile Ala Asn Met .Phe Gly Ser Ser Glu Lys Glu Thr Ile Glu Leu
                                     130
                125
Ser Pro Thr Gly Arg Pro Lys Arg Arg Thr Arg Lys Ser Ile Asn
                140
                                     145
Tyr Ser Lys Ile Asp Asp Phe Pro Asn Glu Leu Glu Lys Leu Ile
                155
                                     160
                                                          165
Ser Gln Ile Gln Pro Glu Val Asp Arg Glu Arg Ala Val Val Glu
                170
                                     175
Val Asn Ile Pro Val Glu Ser Glu Val Asn Leu Lys Leu Gln Asn
                185
                                     190
                                                          195
Ile Met Met Leu Leu Arg Lys Cys Cys Asn His Pro Tyr Leu Ile
                                                          210
                                     205
                200
Glu Tyr Pro Ile Asp Pro Val Thr Gln Glu Phe Lys Ile Asp Glu
                                     220
                215
Glu Leu Val Thr Asn Ser Gly Lys Phe Leu Ile Leu Asp Arg Met
                230
                                     235
                                                          240
Leu Pro Glu Leu Lys Lys Arg Gly His Lys Val Leu Leu Phe Ser
                                     250
                                                          255
                245
Gln Met Thr Ser Met Leu Asp Ile Leu Met Asp Tyr Cys His Leu
                260
                                     265
                                                          270
Arg Asp Phe Asn Phe Ser Arg Leu Met Gly Pro Cys Leu Thr
                                                          Gln
                275
                                     280
                                                          285
Arg Glu Lys Lys Thr Cys Thr Ala Ser Thr Arg Ile Gln Arg Cys
                                     295
                290
Leu Ser Ser
<210> 329
<211> 72
<212> PRT
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<400> 329
Thr Ala Leu Leu Leu Thr Gln Ser Leu Phe Gly Ser Leu Phe Thr.
                                                           15
                                      10
Trp Thr Arg Val Thr Phe Gly Ala Glu Asp Pro Gly Gln Glu Asp
                                                           30
                  20
Ser Phe Arg Arg Val Pro Cys Pro Cys Pro His Ser Val Arg
                  35
                                      40
                                                           45
Arg Ser Thr Tyr Asp Leu Arg Ser Ser Asp Gln Pro Ala Gln Gly
                 50
                                      55
                                                           60
Thr Ser His Glu Phe Gln Ile Gly Phe Pro Thr Ile
                  65
<210> 330
<211> 76
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte ID No: LG:1092522.1.orf2:2000FEB18
<400> 330
Phe Ser Tyr Leu Ser Ser Lys Trp Val Val Lys Gln Gln Arg Gln
                                      10
Leu Ala Ile Ser Thr Met His Leu Ala Gln Glu Leu Leu Met Asn
```

Val Gln Cys Ser Gly Gly Ser Arg His Phe Ser Lys Glu Met Arg

```
Thr Leu Lys Met Arg Ser Ile Val Ala Lys Pro Leu Glu Val Asp
                 50
                                      55
                                                            60
Asn Asp Gln Leu Arg Ala Ile Ser Lys Ala Asp Pro Leu Lys Ala
                 65
<210> 331
<211> 74
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte ID No: LG:1093216.1.orf2:2000FEB18
<220>
<221> unsure
<222> 2, 17, 36
<223> unknown or other
<400> 331
Gly Xaa Pro Pro Thr Thr Ser Gly Pro Gln Thr Asn Gln Pro Lys
                                       10
Glu Xaa Leu Met Asn Phe Lys Ser Asp Ser Gln Leu Tyr Glu Asp
                                       25
                                                            30
                  20
Thr Leu Ala Gly Arg Xaa Val Leu Ile Lys Asn Leu Thr Pro Gln
                  35
                                       40
                                                            45
Thr Leu Gln Pro Arg Trp Thr Gly Pro Tyr Leu Val Ile Tyr Ser
                                       55
                  50
Thr Pro Thr Ala Val Arg Leu Gln Asp Pro Pro His Trp Val
                  65
<210> 332
<211> 67
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte ID No: LI:270318.3.orf3:2000FEB01
Leu Ser Phe Lys Arg Asp Ser Trp Glu Tyr Gly His Pro Ala Pro
                                       10
                                                            15
Arg Cys Gly Asn Glu Ser Ser Arg Ser Gly Glu Ala Ala Leu Ala
                                       25
                                                            30
                  20
Asp Val Gln Leu Ala Ala Pro Val Ser Asn Gln Leu His Pro Asp
                                                            45
                                       40
                  35
Gly Val Glu Asp Arg Gly Val Gly Gly Leu Leu Arg Ser Tyr Thr
                  50
Thr Gln Leu Thr Met Asn Ile
                  65
<210> 333
<211> 192
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte ID No: LI:335671.2.orf2:2000FEB01
Trp Val His Val Leu Leu Arg Glu Arg Lys Lys His Ala Gln Leu
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10
Gln His Gly Ser Arg Gly Val Tyr Leu Leu Val Ser Thr Arg Ala
                 20
                                      25
                                                           30
Gly Gly Leu Gly Ile Asn Leu Thr Ala Ala Asp Thr Val Ile Ile
                                                           45
                 35
                                      40
Tyr Asp Ser Asp Trp Asn Pro Gln Ser Asp Leu Gln Ala Gln Asp
                                      55
                                                           60
                 50
Arg Cys His Arg Ile Gly Gln Thr Lys Pro Val Val Val Tyr Arg
                                      70
                 65
Leu Val Thr Ala Asn Thr Ile Asp Gln Lys Ile Val Glu Arg Ala
                                      85
                 80
Ala Ala Lys Arg Lys Leu Glu Lys Leu Ile Ile His Lys Asn His
                                     100
                 95
Phe Lys Gly Gly Gln Ser Gly Leu Asn Leu Ser Lys Asn Phe Leu
                                     115
                                                          120
                110
Asp Pro Lys Glu Leu Met Glu Leu Leu Lys Ser Arg Asp Tyr Glu
                125
                                     130
                                                          135
Arg Glu Ile Lys Gly Ser Arg Glu Lys Val Ile Ser Asp Lys Asp
                                                          150
                140
                                     145
Leu Glu Leu Leu Asp Arg Ser Asp Leu Ile Asp Gln Met Asn
                155
                                     160
                                                          165
Ala Ser Gly Pro Ile Lys Glu Lys Met Gly Ile Phe Lys Ile Leu
                170
                                     175
                                                          180
Glu Asn Ser Glu Asp Ser Ser Pro Glu Cys Leu Phe
                185
<210> 334
<211> 74
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte ID No: LI:793758.1.orf2:2000FEB01
<220>
<221> unsure
<222> 36
<223> unknown or other
<400> 334
Gly Asp Pro Pro Thr Thr Ser Gly Pro Gln Thr Asn Gln Pro Lys
                                      10
Glu His Leu Met Asn Phe Lys Ser Asp Ser Gln Leu Tyr Glu Asp
                                                           30
                 20
                                      25
Thr Leu Ala Gly Arg Xaa Val Leu Ile Lys Asn Leu Thr Pro Gln
                 35
                                      40
                                                           45
Thr Leu Gln Pro Arg Trp Thr Gly Pro Tyr Leu Val Ile
                                                          Ser
                 50
                                      55
Thr Pro Thr Ala Val Arg Leu Gln Asp Pro Pro His Trp Val
                                      70
                 65
<210> 335
<211> 72
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte ID No: LI:803718.1.orf2:2000FEB01
<220>
<221> unsure
<222> 41
<223> unknown or other
<400> 335
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Thr Ala Leu Leu Thr Gln Ser Leu Phe Gly Ser Leu Phe Thr
                                      10
Trp Thr Arg Val Thr Phe Gly Ala Glu Asp Pro Gly Gln Glu Asp
                                                           30
                                      25
                 20
Ser Phe Arg Arg Val Pro Cys Pro Cys Xaa His Ser Val Arg
                 35
                                      40
Arg Ser Thr Tyr Asp Leu Arg Ser Ser Asp Gln Pro Ala Gln Gly
                 50
Thr Ser His Glu Phe Gln Ile Gly Phe Pro Thr Ile
                 65
<210> 336
<211> 55
<212> PRT
<213> Homo sapiens
<221> misc_feature
<223> Incyte ID No: LI:412179.1.orf2:2000FEB01
<400> 336
Thr Ile Glu Met Met Leu Asp Ile Lys Gln Ile Gln Val Ile Phe
                                      10
                                                           15
Leu Phe Glu Phe Lys Met Gly Arg Lys Ile Ala Glu Thr Thr Arg
                                                           30
                 20
                                      25
Asn Ile Asp Asn Ala Phe Gly Pro Gly Leu Leu Thr Asn Val Gln
                                      40
                 35
Cys Ser Gly Ser Ser Arg Arg Gln Gly Ala
                 50
<210> 337
<211> 107
<212> PRT
<213> Homo sapiens
<220>
<221> misc feature
<223> Incyte ID No: LI:815679.1.orf3:2000FEB01
<400> 337
Leu Arg Tyr Ile Asn Gly Ser Met Ser Ser Leu Tyr Pro Arg Leu
                                      10
Cys His Leu Ser Leu Gln Phe Leu Pro Leu Lys Asn Arg Ser Ile
                                      25
                                                           30
                 20
Phe Leu Gln Ser Leu Met Leu Gly Phe Glu Leu Cys Leu Ala Leu
                                      40
Ala Thr Gly Ile Leu Ile Cys Met Thr Lys Asn Leu Glu Ser Val
                 50
                                      55
                                                           60
Asn Ser Phe Val Leu Ala His Ser Cys Tyr His His Glu Asn Lys 65 70 75
Pro Arg Pro Gly Cys Cys Phe Gln Gln Lys Ile Arg Asp Thr Lys
                                      85
                 80
Asn Lys Val Glu Leu Pro Arg His Ala His Ala Arg Leu Thr Asn
Pro Gln
<210> 338
<211> 147
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte ID No: LI:481361.3.orf3:2000FEB01
<400> 338
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Cys Pro Leu Gln Glu Met Lys Pro Gln Arg Asn Thr Ala Asp Leu
                                     10
Leu Pro Lys Leu Lys Ser Met Ala Leu Ala Asp Arg Ala Val Phe
Glu Lys Gly Met Lys Ala Phe Val Ser Tyr Val Gln Ala Tyr Ala
                                      40
Lys His Glu Cys Asn Leu Ile Phe Arg Leu Lys Asp Leu Asp Phe
                 50
                                      55
                                                          60
Ala Ser Leu Ala Arg Gly Phe Ala Leu Leu Arg Met Pro Lys Met
                 65
                                      70
                                                          75
Pro Glu Leu Arg Gly Lys Gln Phe Pro Asp Phe Val Pro Val Asp
                                                          90
                 80
                                      85
Val Asn Thr Asp Thr Ile Pro Phe Lys Asp Lys Ile Arg Glu Lys
                 95
                                    100
                                                         105
Gln Arg Gln Lys Leu Leu Glu Gln Gln Arg Arg Glu Lys Thr Glu
                                                         120
                                    115
                110
Asn Glu Gly Arg Arg Lys Phe Ile Lys Asn Lys Ala Trp Ser Lys
                                     130
                125
Gln Lys Ala Lys Lys Glu Lys Lys Lys Met Asn
<210> 339
<211> 257
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte ID No: LG:247388.1.orf1:2000MAY19
<400> 339
Gln Asn Ser Val Lys Leu Ala Ile Leu Tyr Leu Met Thr Phe His
                                      10
Leu Gln Ala Met Val Arg Ser Ala Gly Lys Leu Val Leu Ile Asp
                 20
Lys Leu Leu Pro Lys Leu Lys Ala Gly Gly His Lys Val Leu Ile
                                      40
Phe Ser Gln Met Val Arg Cys Leu Asp Ile Leu Glu Asp Tyr Leu
                 50
                                      55
                                                          60
Ile Gln Arg Arg Tyr Leu Tyr Glu Arg Ile Asp Gly Arg Val Arg
                                      70
                 65
Gly Asn Leu Arg Gln Ala Ala Ile Asp Arg Phe Ser Lys
                                                     Pro Asp
                                      85
                 80
Ser Asp Arg Phe Val Phe Leu Leu Cys Thr Arg Ala Gly Gly Leu
                 95
                                     100
Gly Ile Asn Leu Thr Ala Ala Asp Thr Cys Ile Ile Phe Asp Ser
                110
                                     115
                                                         120
Asp Trp Asn Pro Gln Asn Asp Leu Gln Ala Gln Ala Arg Cys His
                125
                                     130
                                                         135
Arg Ile Gly Gln Ser Lys Ala Val Lys Val Tyr Arg Leu Ile Thr
                140
                                     145
                                                         150
Arg Asn Ser Tyr Glu Arg Glu Met Phe Asp Lys Ala Ser Leu Lys
                155
                                                         165
                                     160
Leu Gly Leu Asp Lys Ala Val Leu Gln Ser Met Ser Gly Arg Asp
                170
                                     175
                                                          180
Gly Asn Ile Thr Gly Ile Gln Gln Phe Ser Lys Lys Glu Ile Glu
                                     190
                                                          195
                185
Asp Leu Leu Arg Lys Gly Ala Tyr Ala Ala Ile Met Glu Glu Asp
                                                          210
                                     205
Asp Glu Gly Ser Lys Phe Cys Glu Glu Asp Ile Asp Gln Ile Leu
                215
                                     220
                                                          225
Leu Arg Arg Thr Thr Thr Ile Thr Ile Glu Ser Glu Gly Lys Gly
                                                          240
                230
                                     235
Ser Thr Phe Ala Lys Ala Ser Phe Val Ala Ser Glu Asn Arg
                                                         Thr
                                     250
                245
Asp Ile
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<210> 340
<211> 63
<212> PRT
<213> Homo sapiens
<221> misc_feature
<223> Incyte ID No: LG:255789.10.orf3:2000MAY19
<400> 340
Leu Lys Glu Leu Thr Ile Leu Phe Ser His Phe Pro Ile Gln Met
                                     10
                                                          15
  1
Lys Thr Ala Ser Phe Leu Val Pro Leu Leu Pro Ser Lys Thr Ile
                 20
                                      25
Leu Phe Asp Arg Ala Arg Gly Gln Val Phe Leu Met Leu Leu Arg
                                                          45
                                      40
                 35
Lys Pro Ser Ile Thr Ala His Asp Leu Leu Val Lys Gly Ala Gly
                                                          60
Lys Tyr Lys
<210> 341
<211> 112
<212> PRT
<213> Homo sapiens
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<221> misc_feature
<223> Incyte ID No: LI:787618.1.orf1:2000MAY01
<400> 341
Gly Thr Leu Met Asp Pro Phe Pro Pro Cys Ile Gln Asp Ser Ala
                                      10
                                                          15
Ile Cys Leu Cys Ser Ser Ser Pro Leu Lys Asn Arg Glu Tyr Ile
                                      25
                                                          30
                 20
Ser Pro Ala Pro Asn Val Ala Phe Ser His Met Ser Ser Phe Gly
                                                          45
                 35
                                      40
His Trp Asn Ile Asn Leu His Asp Gln Lys Leu Gly Lys Cys Ala
                 50
                                      55
                                                          60
Phe Ile Cys Ala His Ser Leu Leu Ser Pro Lys Glu Gln Ala
                                      70
                 65
Gln Ala Arg Leu Leu Pro Ala Glu Asp Lys Arg His Gln Glu
                                                          90
                                      85
                 80
Gln Ser Gln Ala Ser Gln Thr Arg Ser Cys Gln Ile Asn Gln Ser
                                     100
                                                         105
                 95
Ser Ala Arg Pro Ile Ala Pro
                 110
<210> 342
<211> 427
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte ID No: LI:331610.2.orf3:2000MAY01
<400> 342
Thr Arg Thr Thr Arg Leu Gly Ser Pro Lys Gly Ser Thr Cys
                                                           15
                                      10
Phe Arg Cys Thr Arg Thr Thr Ser Arg Glu Ser Thr Trp Ala Leu
                                                           30
                                      25
                  20
Cys Ser Pro Arg Ile Pro Thr Trp Ala Lys Asn Gly Thr Val Ser
                                                           45
                  35
                                      40
Tyr Ser Ile Leu Pro Ser His Ile Gly Asp Val Ser Ile Tyr Thr
                  50
                                      55
                                                           60
Tyr Val Ser Val Asn Pro Thr Asn Gly Ala Ile Tyr Ala Leu Arg
```

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65
                                      70
Ser Phe Asn Phe Glu Gln Thr Lys Ala Phe Glu Phe Lys Val Leu
                 80
                                     85
                                                          90
Ala Lys Asp Ser Gly Ala Pro Ala His Leu Glu Ser Asn Ala Thr
                                                          105
Val Arg Val Thr Val Leu Asp Val Asn Asp Asn Ala Pro Val Ile
                110
                                     115
                                                          120
Val Leu Pro Thr Leu Gln Asn Asp Thr Ala Glu Leu Gln Val Pro
                125
                                     130
                                                          135
Arg Asn Ala Gly Leu Gly Tyr Leu Val Ser His Cys Ala Arg Pro
                                                          150
                140
                                     145
Arg Gln Arg Leu Arg Arg Ala Gly Val Ser Pro Thr Lys Ile
                155
                                     160
                                                          165
Val Asp Gly Asn Asp Asp His Leu Phe Glu Ile Asp Pro Ser Ser
                                     175
                                                          180
                170
Gly Glu Ile Arg Thr Leu His Pro Phe Trp Glu Asp Val Thr Pro
                185
                                     190
Val Val Glu Leu Val Val Lys Val Thr Asp His Gly Lys Pro Thr
                200
                                     205
Leu Ser Ala Val Ala Lys Leu Ile Ile Arg Ser Val Ser Gly Ser
                215
                                     220
                                                          225
Leu Pro Glu Gly Val Pro Arg Val Asn Gly Glu Gln His His
                                                          Trp
                230
                                     235
                                                          240
Asp Met Ser Leu Pro Leu Ile Val Thr Leu Ser Thr Ile Ser
                                                          Ile
                                                          255
                245
                                     250
Ile Leu Leu Ala Ala Met Ile Thr Ile Ala Val Lys Cys Lys Arg
                                                          270
                                     265
                260
Glu Asn Lys Glu Ile Arg Thr Tyr Asn Cys Arg Ile Ala Glu Tyr
                                                          285
                275
                                     280
                                                         Ile
Ser His Pro Gln Leu Gly Gly Gly Lys Gly Lys Lys Lys
                290
                                     295
                                                          300
Asn Lys Asn Asp Ile Met Leu Val Gln Ser Glu Val Glu Glu Arg
                305
                                     310
                                                          315
Asn Ala Met Asn Val Met Asn Val Val Ser Ser Pro Ser Leu Ala
                320
                                     325
                                                          330
Thr Ser Pro Met Tyr Phe Asp Tyr Gln Thr Arg Leu Pro Leu Ser
                335
                                     340
Ser Pro Arg Ser Glu Val Met Tyr Leu Lys Pro Ala Ser Asn Asn
                350
                                     355
Leu Thr Val Pro Gln Gly His Ala Gly Cys His Thr Ser Phe Thr
                                     370
                365
Gly Gln Gly Thr Asn Ala Ser Glu Thr Pro Ala Thr Arg Met Ser
                380
                                     385
                                                          390
Ile Ile Gln Thr Asp Asn Phe Pro Ala Glu Pro Asn Tyr Met Gly
                                                          405
                395
                                     400
Ser Arg Gln Gln Phe Val Gln Ser Ile Ser Val Ala Pro Arg Leu
                                     415
                410
Arg Thr Gln Lys Glu Pro Ala
                425
<210> 343
<211> 144
<212> PRT
<213> Homo sapiens
<220>
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<223> Incyte ID No: LG:982697.1.orf2:2000FEB18
<220>
<221> unsure
<222> 52, 56
<223> unknown or other
<400> 343
Gly Ser Ile Glu Gly Lys Cys Gly Val Gly Gly Ser Asn Arg Val
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Thr Ala Gly Ala Leu Pro Asn Gly Thr Ile Arg Ser Gly Pro Leu
                 20
Pro Ser Arg Pro Lys Asp Asp Arg Ser Thr Ser Ser Leu Tyr Ser
                 35
Ala Pro Gly Lys Ala Thr Xaa Thr Gln Leu Xaa Pro Met Ser Ala
                 50
                                      55
                                                          60
Ala Leu Asp Ser Leu Pro Cys Lys Ala Ile Gly Ala Gly Leu Leu
                                      70
                 65
Lys Ala Trp Gly Ala His Pro Leu Tyr Gln Cys Gly Leu Asp Val
                                      85
                 80
Glu His Asp Val Glu Asp Tyr Phe Gly Thr Leu Arg Phe Ser Asp
                 95
                                     100
Phe Pro Thr Gly Phe Trp Ser Cys Val Asp Pro Val Asp Pro Phe
                                                         120
                110
                                     115
Phe Trp Pro Ile Ser Pro Phe Leu Gly Trp Lys His Leu Pro Asn
                125
                                     130
Pro Tyr Thr Pro Ile Val Ser Trp Lys
                140
<210> 344
<211> 97
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte ID No: LG:1080896.1.orf2:2000FEB18
<400> 344
Lys Lys Leu Val Cys Ile Met Leu Phe Tyr Pro Arg Tyr Val Ala
                                      10
Gln Glu Ile Thr Asn Leu Met Arg Val Met Thr His Phe Lys Pro
                 20
                                      25
                                                           30
Pro Met Ile Thr Val Phe Lys Ile Gln Leu Arg Thr Gly Pro Phe
                 35
                                      40
                                                           45
Leu Gly Asp Thr Arg Lys Glu Ile Ala Arg Thr Glu Glu Lys Gly
                                      55
                                                           60
                 50
Val Pro Lys Asn Val Thr Leu Lys Phe Asp Ala Cys Ala Thr Ile
                 65
                                      70
Asp Ser Lys Gln His Gly Ile Gly Cys Gly Ser Leu Asn Trp Lys
                 80
                                      85
Lys Lys Leu His Ser Arg Lys
<210> 345
<211> 75
<212> PRT
<213> Homo sapiens
<221> misc_feature
<223> Incyte ID No: LI:811341.1.orf3:2000FEB01
<400> 345
Gly Leu Phe Gln Cys Ile His Gln Val Thr Glu Val Gly Gln Lys
                                      10
Val Ala Thr Val Leu Leu Phe Tyr Gly Tyr Tyr Lys Cys Thr Gly
                 20
Thr Leu Lys Ile Thr Cys Leu Tyr Asn Val Ile Leu Tyr Lys Val
                 35
                                      40
Cys Ser Pro Gly Ser Asp Gln Pro Asp Val Cys Tyr Asp Pro Ser
                                      55
                                                           60
                 50
Glu Pro Pro Met Thr Thr Val Phe Lys Ile Arg Leu Arg Thr Glu
                                      70
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<210> 346

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<211> 135
<212> PRT
<213> Homo sapiens
<220>
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<223> Incyte ID No: LI:903225.1.orf3:2000FEB01
<400> 346
Asp Pro Phe Gln Lys Met Ala Pro Lys Val Lys Lys Glu Ala Pro
                                      10
Gly Pro Pro Lys Ala Glu Ala Lys Ala Lys Ala Leu Lys Ala Lys
                                                          30
                                      25
                 20
Lys Val Val Leu Lys Gly Val His Gly His Lys Lys Lys Ile
                                                          45
                 35
                                      40
Arg Met Ser Pro Thr Phe Gln Arg Pro Lys Thr Leu Arg Leu Trp
                                      55
                                                          60
                 50
Arg Pro Pro Arg Tyr Pro Arg Lys Thr Thr Pro Arg Arg Asn Lys
                                      70
Leu Asp His Tyr Ala Ile Ile Lys Phe Pro Leu Thr Thr Glu Phe
                                                          90
                 80
                                      85
Ala Met Lys Lys Ile Lys Asp Asn Asn Thr Leu Val Phe Thr Val
                                                          105
                 95
                                     100
Asp Val Lys Ala Asn Lys His Gln Ile Lys Gln Ala Val Lys Lys
                                                          120
                                     115
                110
Leu Cys Asp Ile Asp Gly Ala Lys Val Asn Thr Leu Met Glu Arg
                125
<210> 347
<211> 55
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte ID No: LI:242079.2.orf2:2000FEB01
<400> 347
Ser Ala Arg Ile Glu Ala Trp Glu Pro Pro Thr Arg Phe Gln Ser
                                                          15
                  5
                                      10
Met Cys Gly Lys Ala Trp Met Ser Arg Gln Lys Pro Ala Ala Gly
                                                           30
                 20
                                      25
                    Arg Thr Ser Thr Arg Val Val Trp Arg Gly
Thr Glu Pro Ser Trp
                                      40
                 35
Asn Leu Gly Leu Glu Phe Pro His Ser Phe
                 50
<210> 348
<211> 129
<212> PRT
<213> Homo sapiens
<221> misc_feature
<223> Incyte ID No: LG:979580.1.orf2:2000MAY19
<400> 348
Trp Lys Val Asn Gly Arg Asn Leu Ser Pro Phe Glu Glu Ile Gly
                                                           15
                                      10
Asn Gln Ser His Phe Val Ala Gln Ala Gly Val Gln Trp His Asn
                                                           30
                  20
                                      25
Leu Ala His Cys Asn His His Leu Pro Gly Ser Ser Asp Pro Pro
                                                           45
                 35
                                      40
Thr Ser Thr Ser Gln Val Ala Gly Ser Ala Gly Val Arg His His
                  50
                                      55
Thr Arg Leu Ile Phe Val Phe Leu Val Gln Lys Glu Phe His His
```

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75
Val Asp Gln Ala Gly Leu Lys Leu Leu Thr Ser Ser Asp Trp Pro
                                                           90
                 80
                                      85
Thr Trp Ala Ser Gln Ser Ala Gly Ile Thr Gly Val Ser His
                                                         Cys
                 95
                                     100
                                                          105
Ser Pro Ala Tyr Glu Val Val Phe Ala Val Lys Gln Gln Phe Gly
                110
                                     115
                                                          120
Asn Glu Ala Phe Leu Arg Ser Ser Val
                125
<210> 349
<211> 291
<212> PRT
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<221> misc_feature
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<400> 349
Arg Pro Thr Val Ser Val Ser Cys Ala Ser Ser Arg Pro Gln Phe
                                      10
                                                           15
Leu Ile Thr Val Pro Val Leu Thr Val Ile Asn Tyr Arg Pro His
                                                           30
                 20
                                      25
Asn Met Arg Pro Glu Asp Arg Met Phe His Ile Arg Ala Val Ile
                                                           45
                                      40
                 35
Leu Arg Ala Leu Ser Leu Ala Phe Leu Leu Ser Leu Arg Gly Ala
                 50
                                      55
Gly Ala Ile Lys Ala Asp His Val Ser Thr Tyr Ala Ala Phe Val
                                      70
                                                           75
                 65
Gln Thr His Arg Pro Thr Gly Glu Phe Met Phe Glu Phe Asp Glu
                 80
                                      85
                                                           90
Asp Glu Met Phe Tyr Val Asp Leu Asp Lys Lys Glu Thr Val Trp
                                     100
                                                          105
                  95
His Leu Glu Glu Phe Gly Gln Ala Phe Ser Phe Glu Ala Gln Gly
                                     115
                                                          120
                 110
Gly Leu Ala Asn Ile Ala Ile Leu Asn Asn Asn Leu Asn Thr Leu
                 125
                                     130
Ile Gln Arg Ser Asn His Thr Gln Ala Thr Asn Asp Pro Pro Glu
                 140
                                     145
                                                          150
Val Thr Val Phe Pro Lys Glu Pro Val Glu Leu Gly Gln Pro Asn
                 155
                                     160
                                                          165
Thr Leu Ile Cys His Ile Asp Lys Phe Phe Pro Pro Val Leu Asn
                 170
                                     175
                                                          180
Val Thr Trp Leu Cys Asn Gly Glu Leu Val Thr Glu Gly Val Ala
                                     190
                                                          195
                 185
Glu Ser Leu Phe Leu Pro Arg Thr Asp Tyr
                                         Ser Phe His Lys Phe
                                     205
                 200
                                                          210
His Tyr Leu Thr Phe Val Pro Ser Ala Glu Asp Phe Tyr Asp Cys
                                     220
                                                          225
                 215
Arg Val Glu His Trp Gly Leu Asp Gln Pro Leu Leu Lys His Trp
                 230
                                     235
                                                          240
Glu Ala Gln Glu Pro Ile Gln Met Pro Glu Thr Thr Glu Thr Val
                                     250
                                                          255
                 245
Leu Cys Ala Leu Gly Leu Val Leu Gly Leu Val Gly Ile Ile Val
                                                          270
                                     265
                 260
Gly Ser Val Leu Ile Ile Lys Ser Leu Arg Ser Gly His Asp Pro
                 275
                                     280
Arg Ala Gln Gly Thr Leu
                 290
<210> 350
<211> 517
<212> PRT
<213> Homo sapiens
<220>
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<221> misc\_feature <223> Incyte ID No: LG:337818.2.orf1:2000FEB18

<400> 350 Arg Gly Ala Trp Glu Arg Arg Ser Arg Pro Ala Glu Met Glu Ala Thr Gly Thr Trp Ala Leu Leu Leu Ala Leu Ala Leu Leu Leu Leu Thr Leu Ala Leu Ser Gly Thr Arg Ala Arg Gly His Leu Pro Pro Gly Pro Thr Pro Leu Pro Leu Gly Asn Leu Leu Gln Leu Arg Pro Gly Ala Leu Tyr Ser Gly Leu Met Arg Leu Ser Lys Lys Tyr Gly Pro Val Phe Thr Ile Tyr Leu Gly Pro Trp Arg Pro Val Val Leu Val Gly Gln Glu Ala Val Arg Glu Ala Leu Gly Gly Gln Ala Glu Glu Phe Ser Gly Arg Gly Thr Val Ala Met Leu Glu Gly Thr Phe Asp Gly His Gly Val Phe Phe Ser Asn Gly Glu Arg Trp Arg Gln Leu Arg Lys Phe Thr Met Leu Ala Leu Arg Asp Leu Gly Met Gly Lys Arg Glu Gly Glu Glu Leu Ile Gln Ala Glu Ala Arg Cys Leu Val Glu Thr Phe Gln Gly Thr Glu Gly Arg Pro Phe Asp Pro Ser Leu Leu Leu Ala Gln Ala Thr Ser Asn Val Val Cys Ser Leu Leu Phe Gly Leu Arg Phe Ser Tyr Glu Asp Lys Glu Phe Gln Ala Val Val Arg Ala Ala Gly Gly Thr Leu Leu Gly Val Ser Ser Gln Gly Gly Gln Thr Tyr Glu Met Phe Ser Trp Phe Leu Arg Pro Leu Pro Gly Pro His Lys Gln Leu Leu His His Val Ser Thr Leu Ala Ala Phe Thr Val Arg Gln Val Gln Gln His Gln Gly Asn Leu Asp Ala Ser Gly Pro Ala Arg Asp Leu Val Asp Ala Phe Leu Leu Lys Met Ala Gln Glu Glu Gln Asn Pro Gly Thr Glu Phe Thr Asn Lys Asn Met Leu Met Thr Val Ile Tyr Leu Leu Phe Ala Gly Thr Met Thr Val Ser Thr Thr Val Gly Tyr Thr Leu Leu Leu Leu Met Lys Tyr Pro His Val Gln Lys Trp Val Arg Glu Glu Leu Asn Arg Glu Leu Gly Ala Gly Gln Ala Pro Ser Leu Gly Asp Arg Thr Arg Leu Pro Tyr Thr Asp Ala Val Leu His Glu Ala Gln Arg Leu Leu Ala Leu Val Pro Met Gly Ile Pro Arg Thr Leu Met Arg Thr Thr Arg Phe Arg Gly Tyr Thr Leu Pro Gln Gly Thr Glu Val Phe Pro Leu Leu Gly Ser Ile Leu His Asp Pro Asn Ile Phe Lys His Pro Glu Glu Phe Asn Pro Asp Arg Phe Leu Asp Ala Asp Gly Arg Phe Arg Lys His Glu Ala Phe Leu Pro Phe Ser Leu Gly Lys Arg Val Cys Leu Gly Glu Gly Leu Ala Lys Ala Glu Cys Phe Leu Phe Phe Thr Thr Ile Leu Gln Ala Phe Ser Leu Glu Ser Pro Cys

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475
                470
Pro Pro Asp Thr Leu Ser Leu Lys Pro Thr Val Ser Gly Leu Phe
                485
                                     490
Asn Ile Pro Pro Ala Phe Gln Leu Gln Val Arg Pro Thr Asp Leu
                500
                                     505
                                                          510
His Ser Thr Thr Gln Thr Arg
                515
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Leu Ser Thr Asp Ile Asn Ile Val His Leu Gln Ser Gln Ala His
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Ser Gln Val Thr His Gln Leu Pro Trp Ile Cys Ser Pro His Val
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                                      25
Gly Val Trp Leu Ser Pro Ser Gln Ala Thr Glu Thr Ala Thr His
                 35
                                      40
                                                           45
Val His Ser Ser His Ala Leu Ser Ile His Arg Thr Ser Gln Cys
                                                           60
                 50
                                      55
Pro Cys Pro Trp Cys Leu Ala Gln Gly Thr Ala Cys Pro Leu Arg
                                      70
                 65
Gly His Ala Thr Gln Arg Leu Ser Leu Ser Met Ala Pro Thr His
                                      85
                 80
Ala Pro Ser Leu Gly Tyr Thr Thr Leu Pro Ala Cys Asp His Arg
                                                          105
                 95
                                     100
Cys Pro His Thr Pro Asn His Leu Ser Thr Gln Leu Pro Thr Tyr
                110
                                     115
                                                          120
Asp Ile Val Leu Ala Pro Gln Ser Ile Phe Pro Leu Arg His Ala
                125
                                     130
                                                          135
Ala Pro Thr Glu Ala Gln Ser Pro Ala Thr Ser Ala Thr Ala Ala
                                                          150
                                     145
                140
Leu Ser His Pro Phe Leu Ser Thr Leu Ile Leu Pro Asn Ala Asn
                                                          165
                155
                                     160
Thr Ser Gly Ser Ala Ile Met His Arg Asp Phe Gly His Thr Arg
                                     175
                                                          180
                170
Thr Leu Arg Pro Glu Glu His Leu Pro Asn Pro Asn Thr Cys Leu
                185
                                     190
                                                          195
Cys Asn His Val Glu Ser Gly Pro Cys Cys Pro Ser Thr His Thr
                200
                                     205
                                                          210
Tyr Thr Leu Thr Asp Leu Gln Pro Leu Phe Gly Val Arg Val Pro
                                                          225
                215
                                     220
Thr Arg Pro Ser Gly Arg Gly
                230
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<211> 220
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Gly Lys Val Glu Val Glu Asp Glu Gly Cys Thr Ala Gln Lys Ala
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Pro Val Arg Pro Gly Leu Leu Pro Pro Cys Leu Thr Pro Glu Ile
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                 20
                                      25
Gly Ala Gly Val Pro Ser Ala Gly Cys Pro Leu Cys Pro Ser Met
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Pro Pro Trp Ala Cys Ser Tyr His Thr Ser His Val Pro Met Ile
                 50
                                      55
Pro Leu Leu Gly Pro Arg Ser Ser Phe Ser Arg Lys Trp Ser Ala
Arg Ala Arg Gly Gly Lys Met Ser Pro Tyr Thr Asn Cys
                                                          Tyr
                                                           90
                 80
                                      85
Ala Gln Arg Tyr Tyr Pro Met Pro Glu Glu Pro Phe Cys Thr Glu
                                                          105
                 95
                                     100
Leu Asn Ala Glu Glu Gln Ala Leu Lys Glu Lys Glu Lys Gly Ser
                                                          120
                110
                                     115
Trp Thr Gln Leu Thr His Ala Glu Lys Val Ala Leu Tyr Arg
                                                          Leu
                125
                                     130
                                                          135
Gln Phe Asn Glu Thr Phe Ala Glu Met Asn Arg Arg Ser Asn Glu
                140
                                     145
Trp Lys Thr Val Met Gly Cys Val Phe Phe Phe Ile Gly Phe Ala
                                                          165
                155
                                     160
Ala Leu Val Ile Trp Trp Gln Arg Val Tyr Val Phe Pro Pro Lys
                 170
                                     175
                                                          180
Pro Ile Thr Leu Thr Asp Glu Arg Lys Ala Gln Gln Leu Gln Arg
                185
                                     190
                                                          195
Met Leu Asp Met Lys Val Asn Pro Val Gln Gly Leu Ala Ser Arg
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                200
                                     205
Trp Asp Tyr Glu Lys Lys Gln Trp Lys Lys
<210> 353
<211> 95
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Pro Ile Leu Trp Ser Val Leu Ser Phe Ser Ile Glu Leu Cys Phe
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                                      10
Cys Cys Phe Ile Leu Ser Leu Leu Cys Val Phe Cys Pro Ile Leu
                 20
                                       25
                                                           30
Cys Ser Arg His Gln Glu Pro Arg His Pro Pro Pro Val Thr
                                                          Tyr
                                       40
                 35
Ser Ser Glu Pro Ala Arg Ser Leu Phe Arg Met Ile Thr Trp Arg
                                       55
                                                            60
                 50
Ser Leu Arg Lys Leu Leu Lys Asn Thr Leu Val Pro Ser Leu Ser
                                      70
                                                           75
Gly Leu Gly Pro Phe Arg His Phe Ser Val Ser Met Thr Gln Thr
                 80
                                       85
Met Gln Arg His Phe
                 95
<210> 354
<211> 331
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<220>
<221> unsure
<222> 191
<223> unknown or other
<400> 354
Gly Met Gly Lys Leu Cys Leu Gly Pro Thr Leu Cys Pro Ala Ala
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Ala Ala Glu Glu Ser Arg Asp Ala Glu Pro Arg Arg Glu Leu Leu
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Cys Ser Gly Arg Pro Trp Thr Trp Arg Ala Ala Ala Arg Val Thr
                 35
Thr Met Ile Pro Trp Val Leu Leu Ala Cys Ala Leu Pro Cys Ala
                 50
                                                           60
Ala Asp Pro Leu Leu Gly Ala Phe Ala Arg Arg Asp Phe Arg Lys
                                      70
                 65
Gly Ser Pro Gln Leu Val Cys Ser Leu Pro Gly Pro Gln Gly
                                                           90
                 80
                                      85
Pro Gly Pro Pro Gly Ala Pro Gly Pro Ser Gly Met Met Gly Arg
                                                          105
                 95
                                     100
Met Gly Phe Pro Gly Lys Asp Gly Gln Asp Gly His Asp Gly Asp
                                     115
                                                          120
                110
Arg Gly Asp Ser Gly Glu Glu Gly Pro Pro Gly Arg Thr Gly Asn
                                     130
                125
Arg Gly Lys Pro Gly Pro Lys Gly Lys Ala Gly Ala Ile Gly Arg
                                                          150
                                     145
                140
Ala Gly Pro Arg Gly Pro Lys Gly Val Asn Gly Thr Pro Gly Lys
                155
                                     160
                                                          165
His Gly Thr Pro Gly Lys Lys Gly Pro Lys Gly Lys Lys Gly Glu
                170
                                     175
                                                          180
Pro Gly Leu Pro Gly Pro Cys Ser Cys Gly Xaa Gly His Thr Lys
                                                          195
                185
                                     190
Ser Ala Phe Ser Val Ala Val Thr Lys Ser Tyr Pro Arg Glu Arg
                                                          210
                200
                                     205
Leu Pro Ile Lys Phe Asp Lys Ile Leu Met Asn Glu Gly Gly His
                                     220
                215
Tyr Asn Ala Ser Ser Gly Lys Phe Val Cys Gly Val Pro Gly Ile
                                     235
                                                          240
                230
Tyr Tyr Phe Thr Tyr Asp Ile Thr Leu Ala Asn Lys His Leu Ala
                                                          255
                245
                                     250
Ile Gly Leu Val His Asn Gly Gln Tyr Arg Ile Arg Thr Phe Asp
                                                          270
                260
                                     265
Ala Asn Thr Gly Asn His Asp Val Ala Ser Gly Ser Thr Ile Leu
                                                          285
                275
                                     280
Ala Leu Lys Gln Gly Asp Glu Val Trp Leu Gln Ile Phe Tyr
                                                          Ser
                                     295
                290
Glu Gln Asn Gly Leu Phe Tyr Asp Pro Tyr
                                         Trp Thr Asp Ser Leu
                                     310
                                                          315
                305
Phe Thr Gly Phe Leu Ile Tyr Ala Asp Gln Asp Asp Pro Asn
                                                          Glu
                                     325
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Val
<210> 355
<211> 93
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Asn Val Arg Leu Gln Glu Ile Arg Lys Met Asp Ser Ser Gly Glu
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Ile Tyr Thr Val Gly Ala Glu Val Thr Phe Ser Cys Gln Glu Gly
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                  20
Tyr Gln Leu Met Gly Val Thr Lys Ile Thr Cys Leu Glu Ser Gly
                  35
                                       40
                                                            45
Glu Trp Asn His Leu Ile Pro Tyr Cys Lys Gly Met Phe Ser Lys
                                                            60
                  50
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Phe Thr Thr Phe Leu Met Phe Gly Asn Pro Arg Lys Val Arg
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Arg His Met Lys Cys Tyr Val Val Ser Met Phe Phe Val Phe Glu
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Leu Arg Tyr
<210> 356
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<222> 25, 28, 52, 112
<223> unknown or other
<400> 356
Ser Ser Thr His His Arg Arg Ser Thr Gly Ala Pro Thr Pro Gly
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Leu Pro Pro Pro Pro Pro Ala Thr Arg Xaa Ser Cys Xaa Ala Ala
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Ser Ala Ala Pro Gly Pro Gly Ala Ala Pro Val Gly Ala Pro Thr
                                                           45
                                      40
                 35
Pro Ala Ser Thr Thr Cys Xaa Val Leu Ala Arg Thr Thr Ser Pro
                                      55
                 50
Ser Gly Ser Thr Ala Arg Ser Asp Ala Ala Glu Arg Gly Ser Pro
                 65
Gly Pro Gly Pro Pro Ala Gly Pro Ala Ala Gln Arg His Gly Gly
                 80
                                      85
                                                           90
Arg His Pro His Gly Ser His Leu Asn Pro Gln His Pro Ile Lys
                                                          105
                 95
                                     100
Phe Leu Phe Asn Thr Lys Xaa
                110
<210> 357
<211> 73
<212> PRT
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<223> Incyte ID No: LI:815961.1.orf3:2000FEB01
Cys Val Glu Glu Val Cys Glu Cys Lys Asp Val Glu Val Leu Pro
                                                           15
                                      10
Val Leu Asn Glu Leu Ala Asp Gly Leu Val Pro Leu Val Val Thr
                                                           30
                 20
                                      25
Val Ile Gly Gly Ala Val Trp Val Asp Pro Val Thr Leu Ser Val
                                      40
                 35
Val Ser Gly Gly Met Val Pro Val Gly Val Glu Trp Met Glu Ala
                                      55
                 50
Glu Val Asp Ile Cys Ala Trp Val Gly Val Met Thr Leu
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<210> 358
<211> 239
<212> PRT
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<223> Incyte ID No: LG:120744.1.orf1:2000MAY19
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<221> unsure
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<222> 199 <223> unknown or other <400> 358 Asn Glu Asn Arg Asn Met Ala His Tyr Ile Thr Phe Leu Cys Met Val Leu Val Leu Leu Gln Asn Ser Val Leu Ala Glu Asp Gly Glu Val Arg Ser Ser Cys Arg Thr Ala Pro Thr Asp Leu Val Phe Ile Leu Asp Gly Ser Tyr Ser Val Gly Pro Glu Asn Phe Glu Ile Val Lys Lys Trp Leu Val Asn Ile Thr Lys Asn Phe Asp Ile Gly Pro Lys Phe Ile Gln Val Gly Val Val Gln Tyr Ser Asp Tyr Pro Val Leu Glu Ile Pro Leu Gly Ser Tyr Asp Ser Gly Glu His Leu Thr Ala Ala Val Glu Ser Ile Leu Tyr Leu Gly Gly Asn Thr Lys Thr Gly Lys Ala Ile Gln Phe Ala Leu Asp Tyr Leu Phe Ala Lys Ser Ser Arg Phe Leu Thr Lys Ile Ala Val Val Leu Thr Asp Gly Lys Ser Gln Asp Asp Val Lys Asp Ala Ala Gln Ala Ala Arg Asp Ser Lys Ile Thr Leu Phe Ala Ile Gly Val Gly Ser Glu Thr Glu Asp Ala Glu Leu Arg Ala Ile Ala Asn Lys Pro Ser Ser Thr Tyr Val Phe Tyr Xaa Glu Asp Tyr Ile Ala Ile Ser Lys Ile Arg Glu Val Met Lys Gln Lys Leu Cys Glu Glu Ser Val Cys Pro Thr Arg Ile Pro Val Ala Ala Arg Asp Glu Arg Gly Phe Asp Ile Leu <210> 359 <211> 528 <212> PRT <213> Homo sapiens <221> misc\_feature <223> Incyte ID No: LI:757520.1.orf1:2000MAY01 <400> 359 Ala Ser His Ser Ala Ala Arg Thr Thr Phe Arg Cys Leu Ser Asp Ser Ala Gln Pro His Thr Met Ser Cys Arg Ser Tyr Arg Ile Ser Ser Gly Cys Gly Val Thr Arg Asn Phe Ser Ser Cys Ser Ala Val Ala Pro Lys Thr Gly Asn Arg Cys Cys Ile Ser Ala Ala Pro Tyr Arg Gly Val Ser Cys Tyr Arg Gly Leu Thr Gly Phe Gly Ser Arg Ser Leu Cys Asn Leu Gly Ser Cys Gly Pro Arg Ile Ala Val Gly Gly Phe Arg Ala Gly Ser Cys Gly Arg Ser Phe Gly Tyr Arg Ser Gly Gly Val Cys Gly Pro Ser Pro Pro Cys Ile Thr Thr Val Ser Val Asn Glu Ser Leu Leu Thr Pro Leu Asn Leu Glu Ile Asp Pro 

Asn Ala Gln Cys Val Lys Gln Glu Glu Lys Glu Gln Ile Lys Ser

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Leu Asn Ser Arg Phe Ala Ala Phe Ile Asp Lys Val Arg Phe Leu
                155
                                     160
Glu Gln Gln Asn Lys Leu Leu Glu Thr Lys Trp Gln Phe Tyr Gln
                170
                                     175
                                                          180
Asn Gln Arg Cys Cys Glu Ser Asn Leu Glu Pro Leu Phe Ser Gly
                                     190
                                                          195
                185
Tyr Ile Glu Thr Leu Arg Arg Glu Ala Glu Cys Val Glu Ala Asp
                200
                                     205
                                                          210
Ser Gly Arg Leu Ala Ser Glu Leu Asn His Val Gln Glu Val Leu
                215
                                     220
Glu Gly Tyr Lys Lys Lys Tyr Glu Glu Glu Val Ala Leu Arg Ala
                230
                                     235
                                                          240
Thr Ala Glu Asn Glu Phe Val Val Leu Lys Lys Asp Val Asp Cys
                245
                                     250
                                                          255
Ala Tyr Leu Arg Lys Ser Asp Leu Glu Ala Asn Val Glu Ala Leu
                260
                                     265
                                                          270
Val Glu Glu Ser Ser Phe Leu Arg Arg Leu Tyr Glu Glu Glu Ile
                                                          285
                275
                                     280
Arg Val Leu Gln Ala His Ile Ser Asp Thr Ser Val Ile Val Lys
                                     295
                                                          300
                290
Met Asp Asn Ser Arg Asp Leu Asn Met Asp Cys Ile Ile Ala Glu
                305
                                     310
                                                          315
Ile Lys Ala Gln Tyr Asp Asp Val Ala Ser Arg Ser Arg Ala Glu
                320
                                     325
Ala Glu Ser Trp Tyr Arg Ser Lys Cys Glu Glu Met Lys Ala Thr
                335
                                     340
                                                          345
Val Ile Arg His Gly Glu Thr Leu Arg Arg Thr Lys Glu Glu Ile
                                                          360
                350
                                     355
Asn Glu Leu Asn Arg Met Ile Gln Arg Leu Thr Ala Glu Ile Glu
                                                          375
                365
                                     370
Asn Ala Lys Cys Gln Arg Ala Lys Leu Glu Ala Ala Val Ala Glu
                                                          390
                380
                                     385
Ala Glu Gln Gln Gly Glu Ala Ala Leu Ser Asp Ala Arg Cys Lys
                395
                                     400
Leu Ala Glu Leu Glu Gly Ala Leu Gln Lys Ala Lys Gln Asp Met
                                     415
                                                          420
                 410
Ala Cys Leu Leu Lys Glu Tyr Gln Glu Val Met Asn Ser Lys Leu
                 425
                                     430
                                                          435
Gly Leu Asp Ile Glu Ile Ala Thr Tyr Arg Arg Leu Leu Glu Gly
                                                          450
                 440
                                     445
Glu Glu His Arg Leu Cys Glu Gly Val Gly Ser Val Asn Val Cys
                                     460
                                                          465
                 455
Val Ser Ser Ser Arg Gly Gly Val Ser Cys Gly Gly Leu Ser Tyr
                                                          480
                 470
                                     475
Ser Thr Thr Pro Gly Arg Gln Ile Thr Ser Gly Pro Ser Ala Ile
                 485
                                     490
                                                          495
Gly Gly Ser Ile Thr Val Val Ala Pro Asp Ser Cys Ala Pro Cys
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                 500
                                     505
Gln Pro Arg Ser Ser Phe Ser Cys Gly Ser Ser Arg Ser
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Arg Phe Ala
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<211> 157
<212> PRT
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<223> Incyte ID No: LG:160570.1.orf2:2000FEB18
<220>
<221> unsure
<222> 150
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Val Glu Gly Ala Ala Gly Arg Leu Gly Gly Ser Phe Leu Ser Trp
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Leu Glu Thr Thr Thr Asp Met Gly Cys Phe Cys Ala Val Pro
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                                                           3.0
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Glu Glu Phe Tyr Cys Glu Val Leu Leu Leu Asp Glu Ser Lys Leu
                                      40
                                                           45
                 35
Thr Leu Thr Thr Gln Gln Gly Ile Lys Lys Ser Thr Lys
                                                         Gly
                                      55
                 50
                                                           60
Ser Val Val Leu Asp His Val Phe His His Val Asn Leu Val Glu
                                      70
                 65
Ile Asp Tyr Phe Gly Leu Arg Tyr Cys Asp Arg Ser His Gln Thr
                                      85
                                                           90
                 80
Tyr Trp Leu Asp Pro Ala Lys Thr Leu Ala Glu His Lys Glu Leu
                 95
                                     100
                                                          105
Ile Asn Thr Gly Pro Pro Tyr Thr Leu Tyr Phe Gly Ile Lys Phe
                110
                                     115
                                                          120
Tyr Ala Glu Asp Pro Cys Lys Leu Lys Glu Glu Ile Thr Arg Tyr
                                                          135
                125
                                     130
Ser Ile Asp Phe Val Phe Glu Gln Ile His Ala Leu Arg Ile Xaa
                140
                                                          150
                                     145
Lys Ala Leu Phe Lys Thr Asn
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<210> 361
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<222> 22-23, 56, 65
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Gly Leu Pro Lys Pro Gly Ala Leu Val Gly Asp Arg Ala Ala Pro
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Ala Trp Val Gln Pro Pro Xaa Xaa Gln Val Asn Arg Phe His
                                                         Lys
                                      25
                 20
Ile Arg Asn Arg Ala Leu Leu Thr Asp Gln His Leu Tyr Lys
                                                           45
                 35
                                      40
Leu Asp Pro Asp Arg Gln Tyr Arg Val Met Xaa Ala Val Pro Leu
                 50
Glu Ala Val Thr Xaa
                 65
<210> 362
<211> 517
<212> PRT
<213> Homo sapiens
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<221> misc_feature
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<400> 362
Leu Ala Ala Arg Gly Val Leu Ser Arg Gly Gln Pro Gly Ser Ala
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 1
                                      10
Ala Ala Pro Arg Gln Glu Lys Gln Pro Arg Thr Pro Trp Lys
                                                          Ser
                                                           30
                 20
                                      25
Arg Ser Thr Trp Ala Cys Glu Asn Gly Ala Asn Thr Ser Pro Ala
                 35
Lys Pro His Ser Lys Ala Gly Pro Arg Thr Ala Thr Val Ala Pro
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Ala	Arg	Asn	Leu	Ala 80	Lys	Arg	Arg	Arg	Thr 85	Tyr	Leu	GIA	Gly	Ala 90
Ala	Gly	Arg	Ser	Gln 95	Glu	Pro	Glu	Val	Pro 100	Суѕ	Ala	Ala	Val	Leu 105
Pro	Gly	Lys	Pro	Gly 110	Asp	Arg	Asn	Суѕ	Pro 115	Glu	Phe	Pro	Pro	Pro 120
Asp	Arg	Thr	Leu		Cys	Trp	Ala	Thr		Ala	Ala	Pro	Ala	
Gly	Leu	Суѕ	Gly	Ala 140	Gly	Ser	Glu	Pro		Ile	Ala	Pro	Thr	Ser 150
Cys	Ala	Gly	Asn		Pro	Ser	Arg	Pro	_	Pro	Leu	Leu	Ser	
Leu	Leu	Ala	Ser		Asn	Pro	Cys	Pro		His	Tyr	Leu	His	
Ser	Gly	Ser	His		Thr	Leu	Ala	Pro		Суѕ	Phe	Lys	Ala	
Leu	His	Arg	Lys		Gly	Ser	Gln	Pro		Asp	Met	Ala	Ser	Ala 210
Leu	Thr	Asp	Arg	Thr 215	Ser	Arg	Ala	Pro	Ser 220	Thr	Tyr	Thr	Tyr	Thr 225
Ser	Arg	Pro	Arg	Ala 230	Leu	Pro	Cys	Gln	Arg 235	Ser	Arg	Tyr	Arg	Asp 240
ser	Leu	Thr	Gln	Pro 245	Asp	Glu	Glu	Pro	Met 250	His	Tyr	Gly	Asn	Ile 255
Met	Tyr	Asp	Arg	Arg 260	Val	Ile	Arg	Gly	Asn 265	Thr	Tyr	Ala	Leu	Gln 270
				275					280				Gln	285
Gln	Arg	Glu	Ala	Arg 290	Lys	Arg	Ala	Leu	Ala 295	Arg	Lys	Gln	Ala	Gln 300
			_	305					310			_	Arg	315
His	Val	Asp	Val	Gln 320	Thr	Glu	Leu	Tyr	Leu 325	Glu	Glu	Ile	Ala	Asp 330
Arg	Ile	Ile	Glu	Val 335	Asp	Met	Glu	Cys	Gln 340	Thr	Asp	Ala	Phe	Leu 345
Asp	Arg	Pro	Pro	Thr 350	Pro	Leu	Phe	Ile	Pro 355	Ala	Lys	Thr	Gly	Lys 360
Asp	Val	Ala	Thr	Gln 365	Ile	Leu	Glu	Gly	Glu 370	Leu	Phe	Asp	Phe	Asp 375
Leu	Glu	Val	Lys	Pro 380	Val	Leu	Glu	Val	Leu 385	Val	Gly	Lys	Thr	Ile 390
Glu	Gln	Ser	Leu	Leu 395	Glu	Val	Met	Glu	Glu 400	Glu	Glu	Leu	Ala	Asn 405
				410					415				Ser	420
Arg	Ala	Glu	Val	Gln 425	Arg	Leu	Glu	Glu	Gln 430	Glu	Arg	Arg	His	Arg 435
Glu	Glu	Lys	Glu	Arg 440	Arg	Lys	Lys	Gln	Gln 445		Glu	Ile	Met	His 450
Lys	His	Asn	Glu	Thr 455	Ser	Gln	Lys	Ile	Ala 460		Arg	Ala	Phe	Ala 465
Gln	Arg	Tyr	Leu		Asp	Leu	Leu	Pro	Ser 475		Phe	Gly	Ser	Leu 480
Arg	Asp	Ser	Gly		Phe	Tyr	Asp	Pro		Glu	Arg	Asp	Ile	
Ile	Gly	Phe	Leu		Trp	Leu	Met	Asn			Glu	Lys	Thr	
Glu	Tyr	Ser	Met	Val 515	Gly	Arg			,					

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Met Cys His Ser Gly Gln Met Gly Ser Leu Leu Gly Thr Val Cys
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                   Ser Thr Pro Lys Lys Pro Pro Lys Ser
Ser Glu Ser Val Pro
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Ala Ser Leu Trp Asn Gln Gln Ile Leu His Phe Gly Ala Tyr Lys
<210> 364
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Lys Leu Ala Thr Ser Ser Gly Asp Thr Thr Val Lys Leu Trp Asp
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                                                           30
Leu Cys Thr Gly Asp Cys Ile Leu Thr Phe Glu Gly His Ser Arg
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                 35
                                      40
                    Thr Trp His Ser Cys Gly Asn Phe Val Ala
Ala Val Trp Ser Cys
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                                                           60
                 50
Ser Ser Ser Leu Asp
                    Lys Thr Ser Lys Ile Trp Asp Val Asn Ser
                                      70
                                                           75
                 65
Glu Arg Cys Arg Cys Thr Leu Tyr Gly His Thr Asp Ser Val Asn
                                      85
                                                           90
Ser Ile Glu Phe Phe Pro Phe Ser Asn Thr Leu Leu Thr Ser Ser
                                                          105
                 95
                                     100
Ala Asp Lys Thr Leu Ser Ile Trp Asp Ala Arg Thr Gly Ile Cys
                                                          120
                110
                                     115
Glu Gln Ser Leu Tyr Gly His Met His Ser Ile Asn Asp Ala Ile
                                     130
                                                          135
                125
Phe Asp Pro Arg Gly His Met Ile Ala Ser Cys Asp Ala Cys Gly
                140
                                     145
                                                          150
Val Thr Lys Leu Trp Asp Phe Arg Lys Leu Leu Pro Ile Val Ser
                155
                                     160
                                                          165
Ile Asp Ile Gly Pro Ser Pro Gly Asn Glu Val Asn Phe Asp Ser
                170
                                     175
                                                          180
Ser Gly Arg Val Leu Ala Gln Ala Ser Gly Asn Gly Val Ile His
                                                          195
                185
                                     190
Leu Leu Asp Leu Lys Ser Gly Glu Ile His Lys Leu Met Gly His
                                     205
                                                          210
                200
Glu Asn Glu Ala His Thr Val Val Phe Ser His Asp Gly Glu Ile
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                                     220
                                                          225
Leu Phe Ser Gly Gly Ser Asp Gly Thr Val Arg Thr Trp Ser
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Ala Ala Thr Asn Met Asn Glu Glu Ser Ser Arg Ser His Ala Val

Dh o	Term	<b>T</b> la	(Tibas	230	mb w	ui a	mb ×	T 011	235	7 cm	1727	Laze	Cor	240
	_			Leu 245					250					255
				Lys 260					265					270
Gly	Ser	Glu	Arg	Ala 275	Thr	Lys	Thr	Gly	Ala 280	Ala	Gly	Asp	Arg	Leu 285
Lys	Glu	Gly	Ser	Asn 290	Ile	Asn	Lys	Ser	Leu 295	Thr	Thr	Leu	Gly	Leu 300
Val	Ile	Ser	Ala	Leu 305	Ala	Asp	Gln	Ser	Ala 310	Gly	Lys	Asn	Lys	Asn 315
Lys	Phe	Val	Pro	Tyr 320	Arg	Asp	Ser	Val		Thr	Trp	Leu	Leu	
Asp	Ser	Leu	Gly	Gly 335	Asn	ser	Lys	Thr		Met	Val	Ala	Thr	
Ser	Pro	Ala	Ala	Asp 350	Asn	Tyr	Asp	Glu		Leu	Ser	Thr	Leu	
Tyr	Ala	Asp	Arg	Ala 365	Lys	His	Ile	Val		His	Ala	Val	Val	
Glu	Asp	Pro	Asn	Ala 380	Arg	Ile	Ile	Arg		Leu	Arg	Glu	Glu	
Glu	·Lys	Leu	Arg	Glu 395	Gln	Leu	Thr	Lys		Glu	Ala	Met	Lys	
Pro	Glu	Leu	Lys	Asp 410	Arg	Leu	Glu	Glu		Glu	Lys	Leu	Ile	
Glu	Met	Thr	Val	Thr 425	Trp	Glu	Glu	Lys		Arg	Lys	Thr	Glu	
Ile	Ala	Gln	Glu	Arg 440	Gln	Lys	Gln	Leu		Ser	Leu	Gly	Ile	
Leu	Gln	Ser	Ser	Gly 455	Ile	Lys	Val	Gly		Asp	Lys	Cys	Phe	
Val	Asn	Leu	Asn	Ala 470	Asp	Pro	Ala	Leu		Glu	Leu	Leu	Val	
Tyr	Leu	Lys	Glu	His 485	Thr	Leu	Ile	Gly		Ala	Asn	Ser	Gln	
Ile	Gln	Leu	Cys	Gly 500	Met	Gly	Ile	Leu		Glu	His	Cys	Ile	
Asp	Ile	Thr	ser	Glu 515	Gly	Gln	Val	Met		Thr	Pro	Gln	Lys	Asn 525
Thr	Arg	Thr	Phe	Val 530	Asn	Gly	Ser	Ser		Ser	Ser	Pro	Ile	
Leu	His	His	Gly	Asp 545	Arg	Ile	Leu	$\operatorname{Trp}$		Asn	Asn	His	Phe	
Arg	Leu	Asn	Leu	Pro 560	Lys	Lys	Lys	Lys		Ala	Glu	Arg	Glu	
Glu	Asp	Gln	Asp	Pro 575	Ser	Met	Lys	Asn		Asn	Ser	Ser	Glu	
Leu	Asp	Val	Asp	Gly 590	Asp	Ser	Ser	Ser		Val	Ser	Ser	Glu	
Asn	Phe	Asn	Tyr		Tyr	Ala	Gln	Met		Val	Thr	Met	Lys	Ala 615
Leu	Gly	Ser	Asn		Pro	Met	Gln	Ser		Leu	Asn	Ser	Leu	Glu 630
Gln	Gln	His	Glu	Glu 635	Glu	Lys	Arg	Ser		Leu	Glu	Arg	Gln	
Leu	Met	Tyr	Glu	His 650	Glu	Leu	Glu	Gln		Arg	Arg	Arg	Leu	
Pro	Glu	Lys	Gln	Asn 665	Суз	Arg	Ser	Met		Arg	Phe	Ser	. Phe	
Ser	Pro	Ser	Ala		Gln	Arg	Leu	Arg		Trp	Ala	Glu	Glu	Arg 690
Glu	Ala	Thr	Leu		Asn	Ser	Leu	Met		Leu	Arg	Glu	Gln	Ile 705
Val	Lys	Ala	Asn	Leu	Leu	Val	Arg	Glu	Ala	Asn	Tyr	Ile	Ala	Glu
Glu	Leu	Asp	Lys	710 Arg 725	Thr	Glu	Tyr	Lys	715 Val 730	Thr	Leu	Gln	Ile	720 Pro 735

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Ala Ser Ser Leu Asp Ala Asn Arg Lys Arg Gly Ser Leu Leu Ser
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Glu Pro Ala Ile Gln Val Arg
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Pro Arg Gln His Phe Cys Met Gln Tyr Ser Leu Gln Val Val Thr
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Ala Thr Cys Lys Phe Gly Met Asn Ala Leu Leu Leu Ser Ala Trp
                                                           30
                 20
                                      25
Phe Gly His Leu Arg Ile Leu Gln Ile Leu Val Asn Ser Gly Ala
                                                           45
                                      40
                 35
Lys Ile His Cys Glu Ser Lys Glu Gly Asn Thr Ala Leu His Leu
                 50
                                      55
                                                           60
Ala Ala Gly Arg Gly His Met Ala Val Leu Gln Arg Leu Val Asp
                                                           75
                 65
Ile Gly Leu Asp Leu Glu Glu Gln Asn Ala Glu Gly Leu Thr Ala
                 80
                                      85
                                                           90
Leu His Ser Ala Ala Gly Gly Ser His Pro Asp Cys Val Gln Leu
                                     100
                                                          105
                 95
Leu Leu Arg Ala Gly Ser Thr Val Asn Ala Leu Thr Gln Lys Asn
                 110
                                     115
                                                          120
Leu Ser Cys Leu His Tyr Ala Ala Leu Ser Gly Ser Glu Asp Val
                 125
                                     130
                                                          135
Ser Arg Val Leu Ile His Ala Gly Gly Cys Ala Asn Val Val Asp
                 140
                                     145
                                                          150
His Gln Gly Ala Ser Pro Leu His Leu Ala Val Arg
                155
                                     160
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Ala Leu Val Phe Met Gln Pro Met Val Met Gln Gly Cys Pro Tyr
Thr Leu Pro Arg Cys His Asp Trp Gln Ala Ala Asp Gln Phe His
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                  20
His Ser Ser Ser Leu Arg Ser Thr Cys Pro His Pro Gln Val Arg
                  35
                                      40
                                                           45
Ala Ala Val Thr Ser Pro Ala Pro Pro Gln Asp Gly Ala Gly Val
                                                           60
                  50
                                      55
Pro Cys Leu Ser Leu Lys Leu Leu Asn Gly Ser Val Gly Ala Ser
                                                           75
                                       70
                  65
Gly Pro Leu Glu Pro Pro Ala Met Asn Leu Cys Trp Asn Glu Ile
                  80
                                      85
Lys Lys Lys Ser His Asn Leu Arg Ala Arg Leu Glu Ala Phe Ser
                  95
                                     100
                                                          105
Asp His Ser Gly Lys Leu Gln Leu Pro Leu Gln Glu Ile Ile Asp
                                                          120
                 110
                                     115
Trp Leu Ser Gln Lys Asp Glu Glu Leu Ser Ala Gln Leu Pro Leu
                                      130
Gln Gly Asp Val Ala Leu Val Gln Gln Glu Lys Glu Thr His Ala
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145
                140
Ala Phe Met Glu Glu Val Lys Ser Arg Ala Pro Tyr Ile Tyr Ser
                                     160
                                                          165
                155
Val Leu Glu Ser Ala Gln Ala Phe Leu Ser Gln His Pro Phe Glu
                170
                                     175
Glu Leu Glu Glu Pro His Ser Glu Ser Lys Asp Thr Ser Pro Lys
                                     190
                185
Gln Arg Ile Gln Asn Leu Ser Arg Phe Val Trp Lys Gln Ala Thr
                                     205
                                                          210
                200
Val Ala Ser Glu Leu Trp Glu Lys Leu Thr Ala Arg Cys Val Asp
                215
                                     220
                                                          225
Gln His Arg His Ile Glu Arg Thr Leu Glu Gln Leu Leu Glu Ile
                                                          240
                230
                                     235
Gln Gly Ala Met Glu Glu Leu Ser Thr Thr Leu Ser Gln Ala Glu
                                     250
                                                          255
                245
Gly Val Arg Ala Thr Trp Glu Pro Ile Gly Asp Leu Phe Ile Asp
                                     265
                260
Ser Leu Pro Glu His Ile Gln Ala Ile Lys Leu Phe Lys Glu Glu
                275
                                     280
                                                          285
Phe Ser Pro Met Lys Asp Gly Val Lys Leu Val Asn Asp Leu Ala
                                     295
                                                          300
                 290
His Gln Leu Ala Ile Ser Asp Val His Leu Ser Met Glu Asn Ser
                305
                                     310
                                                          315
Gln Ala Leu Glu Gln Ile Asn Val Arg Trp Lys Gln Leu Gln Ala
                 320
                                     325
                                                          330
Ser Val Asp Glu Arg Leu Lys Gln Leu Gln Asp Ala His Arg Asp
                                     340
                                                          345
                335
Phe Gly Pro Gly Ser Gln His Phe Leu Ser Ser Ser Val Gln Val
                                     355
                                                          360
                350
Pro Trp Glu Arg Ala Ile Ser Pro Asn Lys Val Pro Tyr Tyr Ile
                 365
                                                          375
Asn His Gln Ala Gln Thr Thr Cys Trp Asp His Pro Lys Met Thr
                 380
                                     385
                                                          390
Glu Leu Tyr Gln Thr Leu Ala Asp Leu Asn Asn Ile Lys Phe Ser
                 395
                                     400
                                                          405
Ala Tyr Arg Thr Ala Met Lys Leu Arg Arg Val Gln Lys Ala Leu
                                                          420
                 410
                                     415
Arg Leu Asp Leu Val Thr Leu Thr Thr Ala Leu Glu Ile Phe Asn
                                     430
                                                          435
                 425
Glu His Asp Leu Gln Ala Ser Glu His Val Met Asp Val Val Glu
                 440
                                      445
                                                          450
Val Ile His Cys Leu Thr Ala Leu Tyr Glu Arg Leu Glu Glu
                 455
                                     460
                                                          465
Arg Gly Ile Leu Val Asn Val Pro Leu Cys Val Asp Met Ser Leu
                 470
                                     475
Asn Trp Leu Leu Asn Val Phe Asp Ser Gly Arg Ser Gly Lys Met
                 485
                                     490
                                                          495
Arg Ala Leu Ser Phe Lys Thr Gly Ile Ala Cys Leu Cys Gly Thr
                 500
                                     505
                                                          510
Glu Val Lys Glu Lys Leu Gln Tyr Leu Phe Ser Gln Val Ala Asn
                                      520
                                                          525
                 515
Ser Gly Ser Gln Cys Asp Gln Arg His Leu Gly Val Leu Leu His
                 530
                                     535
                                                          540
Glu Ala Ile Gln Val Pro Arg Gln Leu Gly Glu Val Ala Ala Phe
                 545
                                      550
                                                          555
Gly Gly Ser Asn Val Glu Pro Ser Val Arg Ser Cys Phe Arg Phe
                                                           570
                 560
                                     565
Ser Thr Gly Lys Pro Val Ile Glu Ala Ser Gln Phe Leu Glu Trp
                                                          585
                 575
                                     580
Val Asn Leu Glu Pro Gln Ser Met Val Trp Leu Pro Val Leu His
                                                           600
                 590
                                     595
Arg Val Thr Ile Ala Glu Gln Val Lys His Gln Thr Lys Cys
                                                          Ser
                                                           615
                 605
                                     610
Ile Cys Ser Ala Val Pro His Gln Gly Val Gln Val Pro Glu Ser
                 620
Glu Ala Ile Gln Arg
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<213> Homo sapiens
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Gln Ile Trp Met Lys Ser Thr Gln Ile Ile Gly Leu Gln Tyr Ile
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Ser Leu Pro Leu Gly Met Lys Glu Ile Gly Leu Lys Tyr Lys
                                                         Ara
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                 20
Asp Pro Arg Thr Asn Glu Gly Ile Leu Lys Val Val Lys Ala Leu
                                      40
                                                           45
                 35
Asp Tyr Glu Gln Leu Gln Ser Val Lys Leu Ser Ile Ala Val Lys
                                      55
                                                           60
Asn Lys Ala Glu Phe His Gln Ser Val Ile Ser Arg Tyr Arg
                                                         Val
                                                           75
                 65
                                      70
Gln Ser Thr Pro Val Thr Ile Gln Val Ile Asn Val Arg Glu Gly
                                                           90
                 80
                                      85
Ile Ala Phe Arg Pro Ala Ser Lys Thr Phe Thr Val Gln Lys Gly
                                                          105
                 95
                                     100
Ile Ser Ser Lys Lys Leu Val Asp Tyr Ile Leu Gly Thr Tyr Gln
                110
                                     115
                                                          120
Ala Ile Asp Glu Asp Thr Asn Lys Ala Ala Ser Asn Val Lys Tyr
                                                          135
                125
                                     130
Val Met Gly Arg Asn Asp Gly Gly Tyr Leu Met Ile Asp Ser Lys
                140
                                     145
                                                          150
Thr Ala Glu Ile Lys Phe Val Lys Asn Met Asn Arg Asp Ser Thr
                155
                                     160
                                                          165
Phe Ile Val Asn Lys Thr Ile Thr Ala Glu Val Leu Ala Ile Asp
                170
                                     175
                                                          180
Glu Tyr Thr Gly Lys Thr Ser Thr Gly Thr Val Tyr Val Arg Val
                185
                                     190
                                                          195
Pro Asp Phe Asn Asp Asn Cys Pro Thr Ala Val Leu Glu Lys Asp
                                                          210
                200
                                     205
Ala Val Cys Ser Ser Ser Pro Ser Val Val Val Ser Ala Arg Thr
                215
                                     220
                                                          225
Leu Asn Asn Arg Tyr Thr Gly Pro Tyr Thr Phe Ala Leu Glu Asp
                                                          240
                230
                                     235
Gln Pro Val Lys Leu Pro Ala Val Trp Ser Ile Thr Thr Leu Asn
                                                          255
                245
                                     250
Ala Thr Ser Ala Leu Leu Arg Ala Gln Glu Gln Ile Pro Pro Gly
                260
                                     265
                                                          270
Val Tyr His Ile Ser Leu Val Leu Thr Asp Ser Gln Asn Asn Arg
                275
                                     280
                                                          285
Cys Glu Met Pro Arg Ser Leu Thr Leu Glu Val Cys Gln Cys Asp
                290
                                     295
                                                          300
Asn Arg Gly Ile Cys Gly Thr Ser Tyr Pro Thr Thr Ser Pro Gly
                305
                                     310
                                                          315
Thr Arg Tyr Gly Arg Pro His Ser Gly Arg Leu Gly Pro Ala Ala
                320
                                     325
                                                          330
Ile Gly Leu Leu Leu Gly Leu Leu Leu Leu Leu Ala Pro
                                                          345
                335
                                     340
Leu Leu Leu Thr Cys Asp Cys Gly Ala Gly Ser Thr Gly Gly
                 350
                                     355
                                                          360
Val Thr Gly Gly Phe Ile Pro Val Pro Asp Gly Ser Glu Gly Thr
                                     370
                 365
                                                          375
Ile His Gln Trp Gly Ile Glu Gly Ala His Pro Glu Asp Lys Glu
                 380
                                     385
                                                          390
Ile Thr Asn Ile Cys Val Pro Pro Val Thr Ala Asn Gly Ala Asp
                 395
                                     400
                                                          405
Phe Met Glu Ser Ser Glu Val Cys Thr Asn Thr Tyr Ala Arg Gly
                 410
                                     415
                                                          420
Thr Ala Val Glu Gly Thr Ser Gly Asn Gly Asn Asp His
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425 430

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Ile Leu Thr Arg Asn Met Asp Lys Gln Arg Gln Lys Arg Leu Gln
                                     385
                                                          390
                380
Glu Gln Lys Gln Gln Glu Gly Tyr Asp Gly Gly Pro Asn Leu Arg
                395
                                     400
                                                          405
Thr Lys Val Trp Gln Arg Gly Thr Pro Ser Ser Pro Tyr
                                                         Val
                                                          420
                410
                                     415
Ser Pro Arg His Ser Pro Trp Ser Ser Pro Lys Leu Pro Lys
                                                         Lys
                                                          435
                                     430
                425
Val Gln Thr Val Arg Ser Arg Pro Leu Ser Pro Val Ala Thr Pro
                                     445
                                                          450
                440
Pro Pro Pro Arg Asp Ser Ser Thr Pro Arg Glu Lys Ala His Tyr
                455
                                     460
Gln Lys His Cys Arg Ser His Gln Thr Thr Gly Glu Cys Pro Thr
                                     475
                                                          480
                470
Gly Ile Thr Lys Trp Thr Lys Lys Glu Lys Arg Glu Lys Gly Gln
                485
                                     490
                                                          495
Glu Thr Ala Lys Gln Tyr Ser Lys Gly Asn Lys Lys Phe Ser Glu
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                                     505
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Val Ser Ala Arg Glu Glu Asn Gly Arg Gln Phe Pro Thr Phe
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Pro Thr Glu Ile Ser Ser
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Gln Ser Phe Lys Phe Leu Glu Gln Ile Leu Trp Ser Asn Leu Gln
                                      25
                 2.0
Ile Leu Lys Lys Thr His His Pro Thr His Arg Arg Tyr Asp Phe
                                      40
                 35
Phe Val Ser Arg Phe Ser Ala Met Cys His Ser Cys His Ser Asp
                 50
                                      55
                                                           60
Pro Glu Ile Arg Thr Glu Ile Arg Ile Ala Gly Ile Arg Gly Ile
                                      70
                 65
Gln Gly Val Val Arg Lys Thr Val Asn Asp Glu Leu Arg Ala Thr
                                      85
                                                           90
                 80
Ile Trp Glu Pro Gln His Met Asp Lys Ile Val Pro Ser Leu Leu
                                                          105
                 95
                                     100
Phe Asn Met Gln Lys Ile Glu Glu Val Asp Ser Arg Ile Gly Pro
                110
                                     115
                                                          120
Pro Ser Ser Pro Ser Ala Thr Asp Lys Glu Glu Asn Pro Ala Val
                125
                                     130
                                                          135
Leu Ala Glu Asn Cys Phe Arg Glu Leu Leu Gly Arg Ala Thr Phe
                                     145
                                                          150
                140
Gly Asn Met Asn Asn Ala Val Arg Pro Val Phe Ala His Leu Asp
                155
                                     160
                                                          165
His His Lys Leu Trp Asp Pro Asn Glu Phe Ala Val His Cys Phe
                170
                                     175
                                                          180
Lys Ile Ile Met Tyr Ser Ile Gln Ala Gln Tyr Ser His His Val
                                                          195
                185
                                     190
Ile Gln Glu Ile Leu Gly His Leu Asp Ala Arg Lys Lys Asp Ala
                                                          210
                200
                                     205
Pro Arg Val Arg Ala Gly Ile Ile Gln Val Leu Leu Glu Ala Val
                                                          225
                                     220
                215
Ala Ile Ala Ala Lys Gly Ser Ile Gly Pro Thr Val Leu Glu Val
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                                     235
                230
Phe Asn Thr Leu Leu Lys His Leu Arg Leu Ser Val Glu Phe Glu
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35

40

55

Cys Asn Ala Arg Tyr Ser Ser Pro Arg Pro Lys Leu Leu Glu Asp

45

60

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<212> PRT
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Pro Gly Thr Pro Asn Pro Ala Pro Pro Pro Pro Arg Val His Pro
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Ser Ser Trp Arg Ala Pro Ile Gln Glu Met Ala Val Pro Leu Leu
                 20
                                      25
Thr Lys Lys Ile Val Lys Lys Arg Val Lys Gln Phe Lys Arg Pro
                                      40
                 35
His Leu Asp Arg Tyr Lys Cys Leu Lys Pro Ser Trp Arg Arg Pro
                 50
                                      55
Lys Gly Ile Asp Ser Arg Val Arg Arg Lys Phe Lys Gly Cys
                                                         Thr
                                      70
                                                           75
                 65
Leu Met Pro Asn Ile Gly Tyr Gly Ser Asp Lys Ser Thr Arg His
                 80
                                      85
                                                           90
Tyr Leu Pro Asn Lys Phe Lys Lys Phe Val Val His Asn Val Ser
                                                          105
                 95
                                     100
Glu Leu Glu Leu Met Met His Asn Arg Thr Tyr Cys Ala Glu
                                                          120
                110
                                     115
Ile Ala His Asn Val Ser Thr Lys Lys Arg Lys Glu Ile Val Glu
                125
                                     130
                                                          135
Arg Ala Ala Gln Leu Asp Ile Val Val Thr Asn Lys Leu Ala Arg
                140
Leu Arg Ser Gln Glu Asp Glu
                155
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Ala Ser Cys Ser Arg Arg Arg Glu Ala Leu Gln Arg Thr Ser Val
Asn Met Gly Lys Thr Arg Gly Met Gly Ala Gly Arg Lys Leu Lys
                 20
                                      25
                                                           30
Thr His Arg Arg Asn Gln Arg Trp Ala Asp Lys Ala Tyr Lys Lys
                 35
                                      40
Ser His Leu Gly Asn Glu Trp Lys Lys Pro Phe Ala Gly Ser Ser
                                                           60
                 50
                                      55
His Ala Lys Gly Ile Val Leu Glu Lys Ile Gly Ile Glu Ala Lys
                                      70
                 65
Gln Pro Asn Ser Ala Ile Arg Lys Cys Ala Arg Val Gln Leu Val
                 80
                                      85
                                                           90
Lys Asn Gly Lys Lys Ile Ala Ala Phe Val Pro Asn Asp Gly Cys
                                                          105
                                     100
Leu Asn Tyr Ile Glu Glu Asn Asp Glu Val Leu Ile Ala Gly Phe
                                                          120
                110
                                     115
Gly Arg Lys Gly His Ala Val Gly Asp Ile Pro Gly Val Arg Phe
                125
                                     130
                                                          135
Lys Val Val Lys Val Ser Gly Val Ser Leu Leu Ala Leu Phe
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                                     145
Glu Lys Lys Glu Lys Pro Arg Ser
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<212> PRT
<213> Homo sapiens
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<400> 376
Leu Gly Lys His Arg Arg Pro Pro Pro Pro Lys Asp Gly Arg Arg
                                      10
Arg Gly Tyr Gly Arg Leu Leu Ala Pro Met Ala His Glu Lys
                                                         Lys
                                                           3.0
                 20
                                      25
Leu Ser Asn Pro Met Arg Glu Ile Lys Val Gln Lys Leu Val Leu
                                                           45
                 35
                                      40
Asn Ile Ser Val Gly Glu Ser Gly Asp Arg Leu Thr Arg Ala Ala
                                                           60
                 50
                                      55
Lys Val Leu Glu Gln Leu Ser Gly Gln Thr Pro Val Phe Ser Lys
                                      70
                 65
Ala Arg Tyr Thr Val Arg Ser Phe Gly Ile Arg Arg Asn Glu Lys
                                                           90
                 80
                                      85
Ile Ala Cys Tyr Val Thr Val Arg Gly Glu Lys Ala Met Gln Leu
                                                          105
                 95
                                     100
Leu Glu Ser Gly Leu Lys Val Lys Glu Tyr Glu Leu Leu Arg Arg
                 110
                                     115
                                                          120
Asn Phe Ser Asp Thr Gly Cys Phe Gly Phe Gly Ile Gln Glu His
                                                          135
                125
                                     130
Ile Asp Leu Gly Ile Lys Tyr Asp Pro Ser Thr Gly Ile Tyr Gly
                                                          150
                                     145
                140
Met Asp Phe Tyr Val Val Leu Glu Arg Ala Gly Tyr Arg Val Ala
                                                          165
                155
                                     160
Arg Arg Arg Cys Lys Ser Arg Val Gly Ile Gln His Arg Val
                170
                                     175
                                                          180
Thr Lys Glu Asp Ser Met Lys Trp Phe Gln Val Lys Tyr Glu Gly
                 185
                                     190
                                                          195
Val Ile Leu Lys Gln Gly Ser Gly Leu His Val Pro Pro Leu Thr
                                                          210
                200
                                     205
Cys Gly Gln Asn Ser Ser Leu Val Ser Ser Pro Pro Pro Cys Gln
                215
                                     220
                                                          225
Arg Lys Thr Thr Thr His Leu Ala Arg Leu Phe Trp Val
                230
<210> 377
<211> 102
<212> PRT
<213> Homo sapiens
<220>
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<223> Incyte ID No: LG:246415.1.orf3:2000FEB18
<400> 377
Leu Pro Pro Val Arg Ala Ser Asn Met Met Lys Lys Arg Arg Asn
                                                           15
                                      10
                    Lys Gly Arg Gly His Val Gln Pro Ile
Asn Gly Arg Thr Lys
                                                          Cys
                  20
                                      25
Asp Thr Asn Cys Ala Gln Cys Val Pro Lys Asp Lys Ala Ile Asn
                                      40
                 35
Lys Phe Ile Ile Gly Asn Thr Val Glu Ala Ala Val Arg Asp
                                                           60
                  50
                                      55
Ile Ser Glu Ala Ser Val Phe Asp Ala Tyr Val Leu Pro Lys Leu
                                      70
                                                           75
                  65
Tyr Leu Lys Leu His Tyr Cys Leu Ser Cys Ala Ile His Ser
                                                          Ara
                 80
                                      85
                                                           90
Val Val Arg Asn Arg Ser Cys Glu Ala His Lys Asp
                  95
                                     100
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<210> 378
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<212> PRT
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<400> 378
Val Tyr Trp Cys Arg Glu Leu Ile Leu Lys Arg Gly Gln Ala Lys
                                     10
Val Lys Asn Lys Thr Ile Pro Leu Thr Asp Asn Thr Val Ile Glu
                 20
Glu His Leu Gly Lys Phe Gly Val Ile Cys Leu Glu Asp Leu Ile
                                      40
His Glu Ile Ala Phe Pro Gly Lys His Phe Gln Glu Ile Ser Trp
                                      55
                                                          60
Phe Leu Cys Pro Phe His Leu Ser Val Ala Arg His Ala Thr Lys
                                                          75
                 65
                                     70
Asn Arg Val Gly Phe Leu Lys Glu Met Gly Thr Pro Gly Tyr Arg
                                     85
                 80
Gly Glu Arg Ile Asn Gln Leu Ile Arg Gln Leu Asn
                                     100
<210> 379
<211> 177
<212> PRT
<213> Homo sapiens
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<223> Incyte ID No: LG:1101445.1.orf3:2000FEB18
<400> 379
Gly Thr Met Glu Ala Val Pro Glu Lys Lys Lys Val Ala Ala
                                                          15
                                      10
Ala Pro Gly Thr Leu Lys Lys Lys Val Pro Ala Val Pro Glu
                                      25
                                                          30
                 20
Thr Leu Lys Lys Lys Arg Arg Asn Phe Ala Glu Leu Lys Val Lys
                 35
                                      40
Arg Leu Arg Lys Lys Phe Ala Leu Lys Thr Leu Arg Lys Ala Arg
                                      55
                                                          60
Arg Lys Leu Ile Tyr Glu Lys Ala Lys His Tyr His Lys Glu Tyr
                 65
                                      70
Arg Gln Met Tyr Arg Thr Glu Ile Arg Met Ala Arg Met Ala Arg
                                      85
                                                          90
                 80
Lys Ala Gly Asn Phe Tyr Val Pro Ala Glu Pro Lys Leu Ala Phe
                                     100
                                                         105
                 95
Val Ile Arg Ile Arg Gly Ile Asn Gly Val Ser Pro Lys Val Arg
                                     115
                                                         120
                110
Lys Val Leu Gln Leu Leu Arg Leu Arg Gln Ile Phe Asn Gly Thr
                                     130
                125
Phe Val Lys Leu Asn Lys Ala Ser Val Asn Met Leu Arg Ile Val
                140
                                     145
                                                         150
Glu Pro Tyr Ile Ala Trp Gly Val Pro Gln Pro Glu Val Ser Lys
                155
                                     160
                                                         165
Arg Ala His Leu Gln Thr Arg Leu Trp Gln Asn Gln
                170
                                     175
<210> 380
<211> 86
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
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<223> Incyte ID No: LG:452134.1.orf2:2000FEB18
<400> 380
Leu Pro Ser Tyr Ala Pro Glu Ile Val Pro Ala Thr Leu Arg Arg
                                      10
Ser His Ser Glu Thr Gly Arg Pro Gln Pro Lys Gly Leu Lys Gly
                                                           3.0
                 20
Glu Arg Pro Ala Arg Leu Thr Arg Gly Glu Ala Asp Arg Asp Thr
                 35
                                      40
Tyr Arg Gln Ile Ala Val Pro Pro Asp Ala Asp Arg Lys Ala Glu
                 50
                                      55
Ala Glu Ala Gly Ala Gly Ser Glu Thr Glu Phe Gln Phe Arg Gly
                                      70
                 65
Arg Phe Gly Cys Gly Gly Gly Gln Pro Pro Gln
<210> 381
<211> 97
<212> PRT
<213> Homo sapiens
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<221> misc_feature
<223> Incyte ID No: LI:903021.1.orf1:2000FEB01
Arg Asn Phe Thr Glu Leu Lys Ile Lys Arg Leu Arg Asn Lys Phe
                                      10
                                                           15
Ala Gln Lys Met Leu Leu Lys Ala Arg Arg Lys Leu Ile Tyr Glu
                                                           30
                 20
                                      25
Lys Ala Lys His Tyr His Lys Glu Tyr Met Gln Met Tyr Arg Thr
                 35
                                      40
Glu Ile Gln Ile Ser Arg Ile Ala Arg Lys Ala Gly Asn Phe Tyr
                                                            60
                 50
Val Ser Ala Glu Pro Lys Leu Ala Phe Val Ile Arg Ile Gly Gly
                 65
                                      70
Tyr Gln Leu Gly Glu Pro Lys Gly Leu Lys Gly Val Ala Thr Ser
                 80
                                       85
Leu Pro Ser Ser Asn Leu Gln
<210> 382
<211> 82
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte ID No: LI:246422.1.orf1:2000FEB01
<400> 382
Lys Lys Arg Lys Gln Val Pro Lys Phe Thr Leu Asp Arg Thr His
                                       10
                                                            15
Pro Val Glu Asp Gly Ile Met Asp Ala Ala Asn Phe Glu Gln Phe
                                                            30
                  20
                                       25
Phe Gln Glu Arg Ile Lys Met Asn Gly Lys Ala Gly Asn Phe Gl_{	extstyle Y}
                  35
                                       40
Gly Gly Val Val Thr His Arg Arg Glu Gln Glu Gln Asp Gln Arg
                                       55
                  50
Asp Ile Gln Ala Ala Leu Phe Gln Gln Val Phe Glu Ile Ser His
                  65
Gln Lys Ile Ser Glu Glu Glu
                  80
<210> 383
<211> 180
<212> PRT
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<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte ID No: LG:449404.1.orf1:2000MAY19
<400> 383
Ser Gln Pro Lys Ser Cys Leu Arg Ser Gly His Pro Ser Leu His
                                      10
Ala Thr Met Ser Arg Arg Lys Thr Arg Glu Pro Lys Glu Glu Asn
                 20
Val Thr Leu Gly Pro Thr Val Arg Glu Gly Glu Tyr Val Phe Gly
                                      40
                                                          45
                 35
Val Ala His Ile Phe Ala Ser Phe Asn Asp Thr Phe Ile His Ile
                                      55
                                                          60
                 50
Thr Asp Leu Ser Gly Arg Glu Thr Leu Val Arg Ile Thr Gly Gly
                 65
                                      70
                                                          75
Met Lys Val Lys Ala Asp Arg Asp Glu Ser Ser Pro Tyr Ala Ala
                                      85
                 80
Met Leu Ala Ala Gln Asp Val Ala Gln Arg Cys Lys Glu Leu Gly
                                                         105
                 95
                                     100
Ile Thr Ala Leu His Ile Lys Leu Arg Ala Thr Gly Gly Asn Lys
                110
                                     115
Thr Lys Thr Pro Gly Pro Gly Ala Gln Ser Ala Leu Arg Ala Leu
                125
                                     130
                                                          135
Ala Arg Ser Gly Met Lys Ile Gly Arg Ile Glu Asp Val Thr Pro
                140
                                     145
                                                         150
Val Pro Thr Asp Ser Thr Arg Arg Lys Gly Gly Arg Arg Gly Lys
                155
                                     160
                                                          165
Glu Asp Cys Arg Arg His His Tyr Cys Val Pro Phe Ala Gly Ser
                                                          180
                170
                                     175
<210> 384
<211> 118
<212> PRT
<213> Homo sapiens
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<221> misc_feature
<223> Incyte ID No: LG:449413.1.orf3:2000MAY19
<400> 384
Pro Arg Cys Phe Arg Leu Pro Gln Arg Arg Pro Ser Gln Pro
Val Pro Ser Ser Ala Thr Met Gly Lys Thr Arg Gly Met Gly Ala
                 20
                                      25
                                                           3.0
Gly Arg Lys Leu Lys Thr His Arg Arg Asn Gln Arg Trp Ala Asp
                 35
                                      40
                                                           45
Lys Ala Tyr Lys Lys Ser His Leu Gly Asn Glu Trp Lys Lys Pro
                                                           60
                 50
                                      55
Phe Ala Gly Ser Ser His Ala Lys Gly Ile Val Leu Glu Lys Ile
                                      70
                 65
Gly Ile Glu Ala Lys Gln Pro Asn Ser Ala Ile Arg Lys Cys Ala
                                      85
Arg Val Gln Leu Val Lys Asn Gly Lys Lys Ile Ala Ala Phe Val
                 95
                                     100
Pro Asn Asp Gly Cys Leu Asn Tyr Ile Glu Glu Asn Val
                110
                                     115
<210> 385
<211> 164
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
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<223> Incyte ID No: LG:450105.1.orf2:2000MAY19 Pro Ser Ala Ala Arg Pro Pro Tyr Ser Arg Tyr Arg Ala Arg Arg His His Leu Arg Arg Ser Asn Met Gly Lys Thr Arg Gly Met Gly 20 25 30 Ala Gly Arg Lys Leu Lys Thr His Arg Arg Asn Gln Arg Trp Ala 45 35 40 Asp Lys Ala Tyr Lys Lys Ser His Leu Gly Asn Glu Trp Lys Lys 60 50 Pro Phe Ala Gly Ser Ser His Ala Lys Gly Ile Val Leu Glu Lys 75 65 Ile Gly Ile Glu Ala Lys Gln Pro Asn Ser Ala Ile Arg Lys Cys 90 80 85 Ala Arg Val Gln Leu Val Lys Asn Gly Lys Lys Ile Ala Ala Phe 100 105 95 Val Pro Asn Asp Gly Cys Leu Asn Tyr Ile Glu Glu Asn Asp Glu 115 120 110 Val Leu Ile Ala Gly Phe Gly Arg Lys Gly His Ala Val Gly Asp 125 130 135 Ile Pro Gly Val Arg Phe Lys Val Val Lys Val Ser Gly Val Ser 140 145 150 Leu Leu Ala Leu Phe Lys Glu Lys Glu Lys Pro Arg Ser 1.55 <210> 386 <211> 101 <212> PRT <213> Homo sapiens <220> <221> misc\_feature <223> Incyte ID No: LG:460809.1.orf3:2000MAY19 <400> 386 Ala Trp Val Glu Trp Ala Ser Arg Ser Ala Pro Arg Ala His Arg 10 Glu Ile Gln Lys Phe Ala Met Lys Glu Met Gly Thr Pro Asn Leu 25 20 His Ile Asp Val Arg Leu Asn Lys Ala Leu Trp Ala Lys Gly Ile 40 Arg Asn Val Pro Tyr His Ile His Met Lys Leu Pro Arg Lys Leu 55 60 Asn Glu Asp Glu Asp Ser Pro Asp Lys Leu Tyr Ala Leu Val Pro 65 70 75 Thr Tyr Thr Cys Tyr His Phe His Lys Ser Ile Asp Arg Gln Cys 80 -85 Gly Arg Glu Leu Thr Thr Asp Gly Ser Ile His 95 100 <210> 387 <211> 259 <212> PRT <213> Homo sapiens <220> <221> misc\_feature <223> Incyte ID No: LG:481781.1.orf3:2000MAY19 Arg Ser Val Arg Arg Arg Ser Ser Ser Ser Arg Arg Arg Val 15 Ala Ala Pro His Leu Glu Leu Ala Thr Met Ala Arg Gly Leu Lys 30 20 25 Lys His Leu Lys Arg Leu Asn Ala Pro Lys His Trp Met Leu Asp 35 40

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Lys Leu Gly Gly Ala Phe Ala Pro Lys Pro Ser Ser Gly Pro His
                 50
Lys Ser Arg Glu Cys Leu Pro Leu Ile Leu Ile Ile Arg Asn Arg
                 65
Leu Lys Tyr Ala Leu Thr Tyr Arg Glu Val Ile Ser Ile Leu Met
                                      85
                                                           90
                 80
Gln Arg His Val Leu Val Asp Gly Lys Val Arg Thr Asp Lys Thr
                 95
                                     100
                                                          105
Tyr Pro Ala Gly Phe Met Asp Val Ile Ser Ile Pro Lys Thr Asn
                110
                                     115
                                                          120
Glu Asn Tyr Arg Leu Leu Tyr Asp Thr Lys Gly Arg Phe Arg Leu
                125
                                     130
                                                          135
His Pro Ile Arg Asp Glu Asp Ala Lys Phe Lys Leu Cys Lys Val
                                     145
                                                          150
                140
Arg Ser Val Gln Phe Gly Gln Lys Gly Ile Pro Tyr Leu Asn Thr
                                                          165
                155
                                     1.60
Tyr Asp Gly Arg Thr Ile Arg Tyr Pro Asp Pro Leu Ile Lys Ala
                                     175
                                                          180
                170
Asn Asp Thr Ile Lys Ile Asp Leu Glu Thr Asn Lys Ile Val Asp
                185
                                     190
Phe Ile Lys Phe Asp Val Gly Asn Val Val Met Val Thr Gly Gly
                200
                                     205
                                                          210
Arg Asn Thr Gly Arg Val Gly Val Ile Lys Asn Arg Glu Lys His
                215
                                     220
                                                          225
Lys Gly Ser Phe Glu Thr Ile His Val Glu Asp Ser Trp Ala
                                                          240
                230
                                     235
Gly Ser Pro Pro Val Trp Ala Thr Cys Ser Pro Ser Ala Arg Val
                                     250
                                                          255
                245
Ile Ser Arg Gly
<210> 388
<211> 184
<212> PRT
<213> Homo sapiens
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<223> Incyte ID No: LG:1101153.1.orf2:2000MAY19
<400> 388
Arg Arg Pro Phe Leu Leu Arg Ser Phe Ala Ala Met Val
                                      10
Lys Tyr Ser Gln Glu Pro Gly Asn Pro Thr Lys Ser Ala Lys Ala
                                      25
                 20
Met Gly Arg Asp Leu Arg Val His Phe Lys Asn Thr Arg Glu Thr
                 35
                                      40
                                                           45
Ala Phe Ala Leu Arg Lys Leu Pro Leu Thr Lys Ala Lys Arg Tyr
                                      55
                                                           60
Leu Glu Asp Val Ile Ala His Lys Gln Ala Ile Pro Phe Arg Arg
                 65
                                      70
                                                           75
Tyr Cys Gly Gly Val Gly Arg Thr Ala Gln Ala Lys Ser Arg His
                 80
                                      85
Ser Asn Gly Gln Gly Arg Trp Pro Val Lys Ser Ala Arg Phe Ile
                                     100
                                                          105
                 95
Leu Asp Leu Leu Lys Asn Ala Glu Ser Asn Ala Asp Val Lys Gly
                                                          120
                                     115
                 110
Leu Asp Val Asp Asn Leu Tyr Val Ser His Ile Gln Val Asn Gln
                                                          135
                 125
                                     130
Ala Gln Lys Gln Arg Arg Arg Thr Tyr Arg Ala His Gly Arg Ile
                 140
                                     145
                                                          150
Asn Pro Tyr Met Ser Ser Pro Cys His Ile Glu Leu Ile Leu Ser
                                                          165
                 155
                                     160
Glu Lys Glu Glu Pro Val Lys Lys Glu Ala Asp Asn Ile Val Ala
```

Ala Arg Lys Gln

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<210> 389
<211> 152
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte ID No: LI:257695.20.orf2:2000MAY01
<400> 389
Ala Pro Arg Gly Glu Gly Cys Leu Val His Ala Ser Glu Pro Cys
                                      10
Arg Pro Arg Ala Arg Cys Ser Leu Cys Arg Ser Ser Asp Ala Arg
                 20
                                                           30
Arg Gln Arg Gln Leu Trp Ala His Cys Lys Arg Gly Asn Gly Leu
                                      40
                 35
Ile Lys Val Asn Gly Arg Pro Leu Glu Met Ile Glu Pro Arg Thr
                                      55
Leu Gln Tyr Lys Leu Leu Glu Pro Val Leu Leu Gly Lys Glu
                 65
                                      70
                                                           75
Arg Phe Ala Gly Val Asp Ile Arg Val Arg Val Lys Gly Gly Gly
                                                           90
                 80
                                      85
His Val Pro Gln Ile Tyr Gly Glu Ser Gln Glu Leu Gly Ala Trp
                                                          105
                 95
                                     100
Arg Arg Trp Leu Trp Glu Gly Gly Leu His Ser Ala Pro Val Pro
                110
                                                          120
                                     115
Phe Asn Cys Val Ser Phe Ser Gln Leu Ser Val Ser Pro Ser Pro
                125
                                     130
                                                          135
Lys Pro Trp Trp Pro Ile Thr Arg Asn Val Ser Glu His Gly Ser
                140
                                     145
                                                          150
Phe Pro
<210> 390
<211> 158
<212> PRT
<213> Homo sapiens
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<223> Incyte ID No: LI:455771.1.orf3:2000MAY01
<400> 390
Ala Ser Cys Ser Arg Arg Arg Glu Ala Leu Gln Arg Thr Ser Val
                                                           15
Asn Met Gly Lys Thr Arg Gly Met Gly Ala Gly Arg Lys Leu Lys
                                                           30
                 20
                                      25
Thr His Arg Arg Asn Gln Arg Trp Ala Asp Lys Ala Tyr Lys Lys
                                                           45
                 35
                                      40
Ser His Leu Gly Asn Glu Trp Lys Lys Pro Phe Ala Gly Ser Ser
                                      55
                 50
                                                           60
His Ala Lys Gly Ile Val Leu Glu Lys Ile Gly Ile Glu Ala Lys
                                      70
                                                           75
                 65
Gln Pro Asn Ser Ala Ile Arg Lys Cys Ala Arg Val Gln Leu Val
                 80
                                      85
                                                           90
Lys Asn Gly Lys Lys Ile Ala Ala Phe Val Pro Asn Asp Gly Cys
                 95
                                     100
                                                          105
Leu Asn Tyr Ile Glu Glu Asn Asp Glu Val Leu Ile Ala Gly Phe
                 110
                                     115
                                                          120
Gly Arg Lys Gly His Ala Val Gly Asp Ile Pro Gly Val Arg Phe
                                     130
                125
                                                          135
Lys Val Val Lys Val Ser Gly Val Ser Leu Leu Ala Leu Phe
                140
                                     145
Glu Lys Lys Glu Lys Pro Arg Ser
                155
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<210> 391

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<211> 94
<212> PRT
<213> Homo sapiens
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<223> Incyte ID No: LI:274551.1.orf1:2000MAY01
<400> 391
Pro Phe Thr Val Thr Gln Leu Gln Pro Thr Thr Leu Gln Ser Phe
                                      10
                                                           15
Pro Cys Leu Ser Val Leu Gln Arg Leu Ser His Val Ser Gly Phe
                                                          3.0
                 20
                                      25
Leu Arg Ser Ser Thr Leu Ile Gly Leu Ile Trp Cys Ser Ala Gln
                                                           45
                                      40
                 35
Arg Ala Thr Pro Ser Leu Thr Tyr Ile Gly Ser Ser His Leu Asp
                                      55
                                                           60
                 50
Ala Ser Thr Gln Arg Trp Ala Trp Pro Leu Ser Lys Ala Leu Ala
                                      70
                 65
Ala Leu Gln Val Pro Pro Ala Arg Pro Ser Trp Leu Arg Ala Val
                                      85
Phe Ser Leu Leu
<210> 392
<211> 83
<212> PRT
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<223> Incyte ID No: LI:035973.1.orf3:2000MAY01
<400> 392
Gly Cys Leu Ala Gly Ile Arg Lys Asp Asn Lys Met Lys Gly Thr
                                      10
Ser Pro Phe Gly Lys Cys Arg Asp Met Ile His Lys Leu Cys Cys
                                                           30
Leu Cys Gly Ser Lys Ala Tyr His Leu Gln Lys Ser Thr Cys Gly
                 35
                                      40
                                                           45
Lys Cys Gly Ser Pro Ala Lys Arg Lys Arg Lys Cys Asn Trp Thr
                                                           60
                 50
                                      55
Ala Thr Ala Lys Arg Lys Tyr His Gly Asp Trp Leu Asn Glu Ala
                 65
Pro Lys His Cys Ile Leu Gln Ile
                 80
<210> 393
<211> 174
<212> PRT
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<220>
<221> unsure
<222> 151
<223> unknown or other
<400> 393
Trp Trp Val Cys Asp Gly Cys Leu Cys Phe Arg Thr Thr Pro Ala
                                      10
                                                           15
Val Leu Phe Trp Gln Trp Ile Asn Gln Ser Phe Asn Ala Val Val
                                      25
                                                           3.0
Asn Tyr Thr Asn Arg Ser Gly Asp Ala Pro Leu Thr Val Asn Glu
```

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45
                                      40
Leu Gly Thr Ala Tyr Val Ser Ala Thr Thr Gly Ala Val Ala Thr
                                      55
                                                           60
                 50
Ala Leu Gly Leu Asn Ala Leu Thr Lys His Val Ser Pro Leu Ile
                                      70
                 65
Gly Arg Phe Val Pro Phe Ala Ala Val Ala Ala Ala Asn Cys Ile
                                      85
                 80
Asn Ile Pro Leu Met Arg Gln Arg Glu Leu Lys Val Gly Ile Pro
                 95
                                     100
                                                          105
Val Thr Asp Glu Asn Gly Asn Arg Leu Gly Glu Ser Ala Asn Ala
                110
                                     115
                                                          120
Ala Lys Gln Ala Ile Thr Gln Val Val Val Ser Arg Ile Leu Met
                125
                                     130
                                                          135
Ala Ala Pro Gly Met Gly Ile Pro Pro Phe Ile Met Asn Thr Leu
                                                          150
                140
                                     145
Xaa Lys Lys Ala Phe Leu Lys Arg Phe Pro Met Asp Glu Cys Thr
                155
                                     160
His Ser Ser Trp Val Ser Trp Ile Leu
                170
<210> 394
<211> 183
<212> PRT
<213> Homo sapiens
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<400> 394
Gln Gly Pro Arg Val Leu Leu Ala Met Pro Tyr Leu Pro Asn Ser
                                                           15
                                      10
Ala Gly Tyr His His Leu Cys Gly His Arg Pro Gly Arg Leu Arg
                                                           30
                                      25
                 20
Asp Ser Glu Glu Leu Val Ala Ser Ala Val Gln Pro Arg Leu Gly
                                                           45
                 35
                                      40
Arg Pro Arg His Pro Arg Ala Pro Gly Leu Arg Tyr His Ile Gln
                                                           60
                                      55
His Leu Arg Pro Asp Ser Gln Leu Pro Ala Gly Pro Gly Pro Asp
                 65
                                      70
                                                           75
Pro His Ala Gly Thr Ser Leu His Arg Gly Trp Pro Pro Ala Val
                                                           90
                                      85
                 80
His Ala Gly Ser Ala Thr Ser His Pro Val Pro Gly Gly His Ala
                 95
                                     100
                                                          105
Gly Pro Leu Pro Gly Asp Arg Pro Gln Leu His Glu Gly Tyr Ser
                110
                                     115
                                                          120
Ser Cys Glu His Leu Leu Cys Gly Leu Arg Glu His Glu Ala Gly
                                     130
                                                          135
                125
Leu Gly Gly His Val Gln Val Arg Asp Pro Glu Pro Val Pro Pro
                140
                                     145
                                                          150
Ile Pro His Pro Pro His Leu Ser His Trp Arg Leu Met Ile Gln
                155
                                     160
                                                          165
Pro Gln Asp Pro Tyr Ser Leu Ala Thr Arg Ser Gln Tyr Pro
                                                         Asp
                                                          180
                170
                                     175
Pro Gly Ser
<210> 395
<211> 399
<212> PRT
<213> Homo sapiens
<220>
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<223> Incyte ID No: LI:034583.1.orf1:2000FEB01
<400> 395
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Val Gln His Ala Thr Val Ile Pro Glu Thr Met Ala Gly Thr Gln
                                      10
Gln Leu Ala Asp Trp Arg Asn Thr His Ala His Gly Ser His Tyr
                                                           30
                                      25
Asn Pro Ile Met Gln Gln Pro Ala Leu Leu Thr Gly His Val Thr
                 35
                                      40
Leu Pro Ala Ala Gln Pro Leu Asn Val Gly Val Ala His Val Met
                 50
                                      55
                                                           60
Arg Gln Gln Pro Thr Ser Thr Thr Ser Ser Arg Lys Ser Lys Gln
                 65
                                      70
His Gln Ser Ser Val Arg Asn Val Ser Thr Cys Glu Val Ser Ser
                                                           90
                 80
                                      85
Ser Gln Ala Ile Ser Ser Pro Gln Arg Ser Lys Arg Val Lys Glu
                 95
                                     100
                                                          105
Asn Thr Pro Pro Arg Cys Ala Met Val His Ser Ser Pro Ala Cys
                                                          120
                110
                                     115
Ser Thr Ser Val Thr Cys Gly Trp Gly Asp Val Ala Ser Ser Thr
                125
                                     130
Thr Arg Glu Arg Gln Arg Gln Thr Ile Val Ile Pro Asp Thr Pro
                                     145
                                                          150
                140
Ser Pro Thr Val Ser Val Ile Thr Ile Ser Ser Asp Thr Asp Glu
                155
                                     160
                                                          165
Glu Glu Glu Gln Lys His Ala Pro Thr Ser Thr Val Ser Lys Gln
                170
                                     175
                                                          180
Arg Lys Asn Val Ile Ser Cys Val Thr Val His Asp Ser Pro Tyr
                                     190
                                                          195
                185
Ser Asp Ser Ser Ser Asn Thr Ser Pro Tyr
                                         Ser Val Gln Gln Arg
                200
                                     205
                                                          210
Ala Gly His Asn Asn Ala Asn Ala Phe Asp Thr Lys Gly Ser Leu
                                     220
                                                          225
                215
Glu Asn His Cys Thr Gly Asn Pro Arg Thr Ile Ile Val Pro Pro
                230
                                     235
                                                          240
Leu Lys Thr Gln Ala Ser Glu Val Leu Val Glu Cys Asp Ser Leu
                                                          255
                245
                                     250
Val Pro Val Asn Thr Ser His His Ser Ser Ser Tyr Lys Ser Lys
                                                          270
                260
                                     265
Ser Ser Ser Asn Val Thr Ser Thr Ser Gly His Ser Ser Gly Ser
                                     280
                                                          285
                275
Ser Ser Gly Ala I1e Thr Tyr Arg Gln Gln Arg Pro Gly Pro His
                290
                                     295
                                                          300
Phe Gln Gln Gln Pro Leu Asn Leu Ser Gln Ala Gln Gln His
                305
                                     310
                                                          315
Ile Thr Thr Asp Arg Thr Gly Ser His Arg Arg Gln Gln Ala Tyr
                320
                                     325
                                                          330
Ile Thr Pro Thr Met Ala Gln Ala Pro Tyr Ser Phe Pro His Asn
                                     340
                                                          345
                335
Ser Pro Ser His Gly Thr Val His Pro His Leu Ala Ala Ala Ala
                350
                                     355
                                                          360
Ala Ala Ala His Leu Pro Thr Gln Pro His Leu Tyr Thr Tyr Thr
                                     370
                                                          375
                365
Ala Pro Ala Ala Leu Gly Ser Thr Gly Thr Val Ala His Leu Val
                380
                                                          390
Ala Ser Gln Gly Ser Ala Arg His Thr
                395
<210> 396
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<223> Incyte ID No: LI:3333307.2.orf1:2000FEB01
<220>
<221> unsure
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<222> 286

<223> unknown or other

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<400> 396
Thr Gln Phe Ser Asp Asn Asn Glu Lys Leu Ser Pro Lys Pro
                                      10
Gly Thr Gly Glu Pro Val Leu Ser Leu His Tyr Ser Thr Glu Gly
                 20
                                      25
Thr Thr Thr Ser Thr Ile Lys Leu Asn Phe Thr Asp Glu Trp Ser
                 35
                                      40
Ser Ile Ala Ser Ser Ser Arg Gly Ile Gly Ser His Cys Lys Ser
                                      55
Glu Gly Gln Glu Glu Ser Phe Val Pro Gln Ser Ser Val Gln Pro
                 65
                                      70
                                                          75
Pro Glu Gly Asp Ser Glu Thr Lys Ala Pro Glu Glu Ser Ser Glu
                                                          90
                 80
                                      85
Asp Val Thr Lys Tyr Gln Glu Gly Val Ser Ala Glu Asn Pro Val
                                                         105
                 95
                                    100
Glu Asn His Ile Asn Ile Thr Gln Ser Asp Lys Phe Thr Ala Lys
                                    115
                                                         120
                110
Pro Leu Asp Ser Asn Ser Gly Glu Arg Asn Asp Leu Asn Leu Asp
                                    130
                                                         135
                125
Arg Ser Cys Gly Val Pro Glu Glu Ser Ala Ser Ser Glu Lys Ala
                140
                                     145
Lys Glu Pro Glu Thr Ser Asp Gln Thr Ser Thr Glu Ser Ala Thr
                155
                                     160
Asn Glu Asn Asn Thr Asn Pro Glu Pro Gln Phe Gln Thr Glu Ala
                170
                                     175
                                                         180
Thr Gly Pro Ser Ala His Glu Glu Thr Ser Thr Arg Asp Ser Ala
                                     190
                                                         195
                185
Leu Gln Asp Thr Asp Asp Ser Asp Asp Pro Val Leu Ile Pro
                                     205
                                                         210
                200
Gly Ala Arg Tyr Arg Ala Gly Pro Gly Asp Arg Phe Asn Ile Arg
                215
                                     220
                                                         225
Gly Thr Thr Ile Gly Asp Arg Ile Met Arg Arg Ser Ala Val Ala
                230
                                     235
                                                         240
Arg Ile Gln Glu Phe Phe Arg Arg Lys Glu Arg Lys Glu Met
                245
                                     250
                                                         255
Glu Glu Leu Asp Thr Leu Asn Ile Arg Arg Pro Leu Val Lys Met
                260
                                     265
                                                         270
Val Tyr Lys Gly His Arg Asn Ser Arg Thr Met Ile Lys Glu Ala
                                     280
                                                         285
                275
Xaa Phe Trp Gly Ala Asn Phe Val Met Ser Gly Ser Asp Cys Gly
                290
His
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<210> 397

<211> 105

<212> PRT <213> Homo sapiens

<220>

<221> misc\_feature

<223> Incyte ID No: LI:814710.2.orf2:2000FEB01

<400> 397

Glu Gly Pro Glu Gly Lys Trp Lys Ser Pro Lys Phe Lys Met Pro
1 5 10 15

Glu Met His Phe Lys Thr Pro Lys Ile Ser Met Pro Asp Ile Asp
20 25 30

Leu Asn Leu Thr Gly Pro Lys Ile Lys Gly Asp Val Asp Val Thr
35 40 45

Gly Pro Lys Val Glu Gly Asp Leu Lys Gly Pro Glu Val Asp Leu
50 55 60

Lys Gly Pro Lys Val Asp Ile Asp Val Pro Asp Val Asn Val Gln
65 70 75

Gly Pro Asp Trp His Leu Lys Met Pro Lys Met Lys Met Pro Lys

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85
                                                          90
                 80
Phe Ser Met Pro Gly Phe Lys Gly Glu Gly Pro Gly Ser Arg Ser
                                                         105
<210> 398
<211> 153
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Trp Tyr Arg Arg Leu Leu Arg Glu Ser Gly Ser Thr Met Asp Ile
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Pro Val Pro Ser Ser Phe Asn Asp Val Gly Gln Asp Trp Arg Leu
                 20
                                                          30
Arg His Phe Val Asp Gln Met Trp Tyr Glu Arg Glu Val Thr Phe
                 35
                                      40
Leu Glu Gln Trp Thr Gln Asp Leu His Thr Arg Val Val Leu Arg
                 50
                                                          60
                                      55
Ile Val Ser Ala His Ser Tyr Ala Ile Val Trp Val Asn Gly Val
                                      70
Asp Ala Leu Glu His Glu Gly Ser Thr Ser Pro Leu Thr Pro Thr
                 80
                                      85
                                                          90
Ser Val Ala Cys Ser Arg Trp Gly Pro Cys Pro Pro Ala Ser Ala
                 95
                                     100
                                                         105
Ser Leu Ser Pro Ser Ala Thr Cys Ser Ser Pro Pro Pro Cys His
                110
                                     115
                                                         120
Gln Gly Ala Ser Ser Thr Trp Pro Thr Pro Pro Arg Gly Tyr His
                125
                                     130
Pro Ala Ser Thr Ala Asp Thr His Leu Pro Val Pro Pro Arg
                                                         Glv
                140
                                     145
                                                         150
Ala Leu His
<210> 399
<211> 161
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Ser Met Leu Ala Ser Gln Gly Val Leu Leu His Pro Tyr Gly Val
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Pro Met Ile Val Pro Ala Ala Pro Tyr Leu Pro Gly Leu Ile Gln
                 20
                                      25
Gly Asn Gin Glu Ala Ala Ala Pro Asp Thr Met Ala Gln Pro
                 35
                                      40
                                                           45
Tyr Ala Ser Ala Gln Phe Ala Pro Pro Gln Asn Gly Ile Pro Ala
                                      55
                 50
                                                           60
Glu Tyr Thr Ala Pro His Pro His Pro Ala Pro Glu Tyr Thr Gly
                                      70
                 65
Gln Thr Thr Val Pro Glu His Thr Leu Asn Leu Tyr Pro Pro Ala
                 80
                                      85
Gln Thr His Ser Glu Gln Ser Pro Ala Val Phe Leu Phe Val Ile
                                     100
                                                          105
                 95
Thr Arg Ala Val Ala Leu Phe Thr Ser Ile Leu Arg Pro Ser Thr
                110
                                     115
                                                         120
Thr Val Pro Cys Asn Phe Ser Leu Ala Leu Ser Ala Ser Ala Leu
                125
                                     130
                                                          135
Phe Ser Lys Val Thr Lys Pro Asn Pro Phe Glu Pro Arg Ser Leu
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150
                140
Lys Ile Ile Ser Thr Ser Lys Ile Leu Pro Asn
                155
<210> 400
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Trp Tyr Arg Arg Leu Leu Arg Glu Ser Gly Ser Thr Met Asp Ile
                                      10
Pro Val Pro Ser Ser Phe Asn Asp Val Gly Gln Asp Trp Arg Leu
                                      25
                 20
Arg His Phe Val Asp Gln Met Trp Tyr Glu Arg Glu Val Thr Phe
                 35
                                      40
Leu Glu Gln Trp Thr Gln Asp Leu His Thr Arg Val Val Leu Arg
                                      55
                                                           60
                 50
Ile Val Ser Ala His Ser Tyr Ala Ile Val Trp Val Asn Gly
                                                         Val
                 65
                                      70
Asp Ala Leu Glu His Glu Gly Ser Thr Ser Pro Leu Thr Pro Thr
                                                           90
                 80
                                      85
Ser Val Ala Cys Ser Arg Trp Gly Pro Cys Pro Pro Ala Ser Ala
                 95
                                     100
                                                          105
Ser Leu Ser Pro Ser Ala Thr Cys Ser Ser Pro Pro Cys His
                110
                                     115
                                                          120
Gln Gly Ala Ser Ser Thr Trp Pro Thr Pro Pro Arg Gly Tyr His
                125
                                     130
                                                          135
Pro Ala Ser Thr Ala Asp Thr His Leu Pro Val Pro Pro Arg Gly
                                     145
Ala Leu His
<210> 401
<211> 135
<212> PRT
<213> Homo sapiens
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<223> Incyte ID No: LI:900264.2.orf3:2000MAY01
Arg Gln Ile Val Ser Val Leu Ser Cys Arg Phe Asp Ser Asn Gln
                                      10
Leu Pro Ala Asn Ala Pro Ile Glu Asp Arg Arg Ser Ala Ala Thr
                 20
                                      25
                                                           30
Cys Leu Gln Thr Arg Gly Leu Leu Cly Val Phe Asp Gly His
                                      40
                 35
                                                           45
Ala Gly Cys Ala Cys Ser Gln Ala Val Ser Glu Arg Leu Phe Tyr
                 50
                                      55
                                                           60
Tyr Ile Ala Val Ser Leu Leu Pro His Glu Thr Leu Leu Glu Ile
                 65
                                      70
Glu Asn Ala Val Glu Ser Gly Arg Ala Leu Leu Pro Ile Leu Gln
                 80
                                      85
Trp His Lys His Pro Asn Asp Tyr Phe Ser Lys Glu Ala Ser Lys
                 95
                                     100
                                                          105
Leu Tyr Phe Asn Ser Leu Arg Thr Tyr Trp Gln Gly Ala Tyr Arg
                110
                                     115
                                                          120
Pro Gln His Trp Val Ser Arg Leu Ile Leu Met Leu Arg Arg Leu
                125
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<212> PRT
<213> Homo sapiens
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<220>
<221> unsure
<222> 28, 39-40
<223> unknown or other
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Arg Gly Ala Gly Thr Ala Ala Leu Pro Ser Arg Leu Leu Pro Ser
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Ser Ala Ala Arg Ser Ser Leu Gly Ile His Leu Leu Xaa Leu Leu
                 20
                                      25
Leu Leu Ile His Ser Phe Pro His Xaa Xaa Leu Leu Gly Phe
                 35
                                      40
Ser Pro Arg Pro Ala Ser Pro Arg Ala Leu Pro Leu Pro Leu Pro
                 50
                                      55
                                                          60
Val Leu Pro Gly Pro Leu Leu Pro Leu Ile His Ser Pro Leu Ser
                 65
                                      70
                                                          75
Leu Leu His Ser Leu Pro Leu Ser Pro Phe Phe Phe Phe His
                                      85
                                                          90
                 80
Pro Pro Ser Leu Thr Pro Pro Pro Phe Pro Cys Leu Leu Ser Asp
                 95
                                     100
                                                         105
Thr Ala Leu Gln Leu Leu Ser Pro Ala Pro Ser Pro Val Arg
                110
                                     115
Thr Asn Gln Gln His Cys Phe Phe Ser
                125
<210> 403
<211> 299
<212> PRT
<213> Homo sapiens
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<223> Incyte ID No: LI:1189543.1.orf1:2000MAY01
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Glu Phe Arg Gln Asn Lys Arg Glu Asn Leu Leu Pro Val Ala Ala
                  5
                                      10
Ala Gly Thr Ala Asn Met Met Ala Ala Ala Pro Ile Gln Gln Asn
                                                          30
                 20
                                      25
Gly Thr His Thr Gly Val Pro Ile Asp Leu Asp Pro Pro Asp Ser
                 35
                                      40
                                                          45
Arg Lys Arg Pro Leu Glu Ala Pro Pro Glu Ala Gly Ser Thr Lys
                 50
                                      55
                                                          60
Arg Thr Asn Thr Gly Glu Asp Gly Gln Tyr Phe Leu Lys Val Leu
                 65
                                      70
                                                          75
Ile Pro Ser Tyr Ala Ala Gly Ser Ile Ile Gly Lys Gly Gly Gln
                 80
                                      85
                                                          90
Thr Ile Val Gln Leu Gln Lys Glu Thr Gly Ala Thr Ile Lys Leu
                 95
                                     100
Ser Lys Leu Ser Lys Ser Lys Asp Phe Tyr Pro Gly Thr Thr Glu
                110
                                     115
                                                         120
Arg Val Cys Leu Ile Gln Gly Thr Val Glu Ala Leu Asn Ala Val
                125
                                     130
                                                         135
His Gly Phe Ile Ala Glu Lys Ile Arg Glu Met Pro Gln Asn Val
                140
                                     145
                                                         150
Ala Lys Thr Glu Pro Val Ser Ile Leu Gln Pro Gln Thr Thr Val
                155
                                     160
                                                         165
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Asn Pro Asp Arg Ile Lys Gln Thr Leu Pro Ser Ser Pro Thr Thr

```
175
Thr Lys Ser Ser Pro Ser Asp Pro Met Thr Thr Ser Arg Ala Asn
                                                          195
                185
                                     190
Gln Val Lys Ile Ile Val Pro Asn Ser Thr Ala Gly Leu Ile Ile
                200
                                                          210
                                     205
Gly Lys Gly Gly Ala Thr Val Lys Ala Val Met Glu Gln Ser
                                                          Gly
                                                          225
                215
                                     220
Ala Trp Val Gln Leu Ser Gln Lys Pro Asp Gly Ile Asn Leu Gln
                230
                                     235
                                                          240
Glu Arg Val Val Thr Val Ser Gly Glu Pro Glu Gln Asn Arg
                                                          Lys
                245
                                     250
                                                          255
Ala Val Glu Leu Ile Ile Gln Lys Ile Gln Glu Asp Pro Gln Ser
                                     265
                                                          270
                260
Gly Ser Cys Leu Asn Ile Ser Tyr Ala Asn Val Thr Gly Pro
                                                          Va1
                275
                                     280
                                                          285
Gly Lys Phe Gln Ser Asn Arg Ile Ser Leu Cys Lys His Cys
                290
                                     295
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<211> 142
<212> PRT
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Gly Gly Gly Ile Leu Arg Arg Ser Gly Ser Ser Ser Ser Ser Ser
                                                           15
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Ser Ser Ser Glu Asp Asp Gly Met Gly Gly Arg Arg Lys Lys
                                      25
                                                           30
Gly Leu Lys Glu Lys Ile Lys Glu Lys Met Pro Gly Gly His Arg
                 35
                                      40
                                                           45
Glu Gly Gln Gly Gln Ala Thr Ala Thr Gly Ala Tyr Gly Gly
                                                          Thr
                 50
                                      55
                                                           60
Gly Tyr Val Ala Gly Pro Thr Thr Gly Gly
                                         Pro His Glu Lys
                                      70
                                                           75
                 65
Gly Val Val Glu Lys Ile Lys Glu Lys Ile Pro Gly Gly His Lys
                 80
                                      85
                                                           90
Asp Tyr Asp Gln His Gln His Thr Thr Ala Ala Thr Gly Gly Gly
                 95
                                     100
                                                          105
Gly Gly Tyr Gly Gly Thr Thr Asp Thr Thr Tyr Gly Thr
                                                          Thr
                110
                                     115
                                                          120
Thr Glu Gly Thr His Glu Lys Lys Gly Phe Met Asp Lys Ile Lys
                125
                                     130
                                                          135
Glu Lys Leu Pro Gly Gln. His
                140
<210> 405
<211> 168
<212> PRT
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Ser Tyr Leu Arg Ser Arg Gly Gln Pro Pro Pro Arg Arg Ser His
 1
                                      10
                                                           15
Ala Leu Arg Ala Arg Arg Leu Ser Ser Val Ser Ala Ser Leu Pro
                  20
                                      25
                                                           30
Leu Pro Ser Arg Leu Thr His Met Ala Ser Ile Ala Gly Ser
                                                          Ser
                 35
                                      40
                                                           45
Ala Leu Ser Phe Ala Arg Pro Val Lys Ala Ile Asn Thr Asn Ser
                 50
                                      55
```

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Leu Ala Phe Ser Pro Ala Arg Lys Gly Asn Thr Phe Leu Arg Leu
                 65
Gln Pro Met Pro Met Arg Ser Val Ser Cys Ala Ala Lys Lys Asp
                                      85
                                                           90
Thr Thr Asp Lys Val Cys Glu Ile Val Lys Lys Gln Leu Ala Leu
                 95
                                     100
                                                          105
Pro Asp His Thr Glu Val Cys Gly Glu Ser Lys Phe Ser Glu Leu
                110
                                     115
                                                          120
Gly Ala Asp Ser Leu Asp Thr Val Glu Ile Val Met Ser Leu Glu
                                                          135
                125
                                     130
Glu His Phe Asp Ile Ser Val Glu Glu Ser Ser Ala Gln Thr Ile
                                     145
                                                          150
                140
Ala Thr Val Glu Asp Ala Ala Asp Leu Ile Asp Lys Leu Val Ala
                155
                                     160
Gly Lys Ala
<210> 406
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Leu Lys Ala Arg Glu Gln Ala Gln Lys Arg Glu Ala Ile Gln Val
                                                           15
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                    Leu Arg Leu Ile Leu Arg Lys Ala Gly Glu
Thr Ser Pro Val Cys
                                                           30
                 20
                                      25
Glu Val Lys Arg Leu Lys Thr Gln Pro Thr Asp Glu Glu Met Leu
                                      40
                 35
Phe Ile Tyr Ser His Phe Lys Gln Ala Thr Val Gly Asp Val Asn
                                      55
                                                           60
                 50
Thr Asp Arg Pro Gly Leu Leu Asp Leu Lys Gly Lys Ala Lys Trp
                 65
                                      70
Asp Ser Trp Asn Lys Leu Lys Gly Thr Ser Lys Glu Asn Ala Met
                 80
                                      85
                                                           90
Lys Thr Tyr Val Glu Lys Val Glu Glu Leu Lys Lys Lys Tyr Gly
                 95
                                     100
Ile Leu Thr Thr Arg Phe Gly Gly Gln Pro His Val
                110
<210> 407
<211> 804
<212> PRT
<213> Homo sapiens
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<223> Incyte ID No: LG:132147.3.orf3:2000FEB18
<400> 407
Ala Ser Ser Ala Ala Gln Asp Asp Tyr Ala Arg His Val Arg Asp
                                                           15
                                      10
Pro Pro Ala Val Ala Val Ala Ala Pro Lys Ala Arg Ala Ser Ala
                                      25
                 2.0
Pro Asp Ala Val Val Thr Pro Pro Arg Leu Ala Val Ala Pro Val
                 35
                                      40
Pro Val Leu Pro Ala Ala Ser Ala Ala Gly Asp Met Ser Asn Pro
                                      55
                                                           60
Gly Ser Arg Arg Asn Gly Pro Val Lys Leu His Leu Thr Val Leu
                                      70
                                                           75
                 65
Cys Ala Lys Asn Leu Val Lys Lys Asp Phe Phe Arg Leu Pro Asp
                                      85
Pro Phe Ala Lys Val Val Val Asp Gly Ser Gly Gln Cys His Ser
```

Thr	Asp	Thr	Val	.95 Lvs	Asn	Thr	Leu	Asp	100 Pro	Lvs	Trp	Asn	Gln	105 His
	Asp	•		110 Ile					115					120
_	_		Lys	125					130					135
			-	140			-	_	145	_		_		150
_	_		Arg	155					160			_	_	165
	_	_	Gln	170		-		-	175		_			180
	-		Val	185	-				190					195
Asp	Arg	Ile	Gly	Thr 200	Gly	Gly	Gln	Val	Val 205	Asp	Cys	Ser	Arg	Leu 210
Phe	Asp	Asn	Asp	Leu 215	Pro	Asp	Gly	Trp	Glu 220	Glu	Arg	Arg	Thr	Ala 225
Ser	Gly	Arg	Ile		Tyr	Leu	Asn	His		Thr	Arg	Thr	Thr	Gln 240
$\mathtt{Trp}$	Glu	Arg	Pro		Arg	Pro	Ala	Ser		Tyr	ser	Ser	Pro	
Arg	Pro	Leu	Ser		Phe	Val	Asp	Glu		Thr	Pro	Ile	Ser	
Thr	Asn	Gly	Ala		Суз	Gly	Gln	Ser		Asp	Pro	Arg	Leu	
Glu	Arg	Arg	Val	Arg	Ser	Gln	Arg	His		Asn	Tyr	Met	Ser	Arg
Thr	His	Leu	His		Pro	Pro	Asp	Leu	Pro	Glu	Gly	Tyr	Glu	
Arg	Thr	Thr	Gln		Gly	Gln	Val	Tyr		Leu	His	Thr	Gln	
Gly	Val	Ser	Thr		His	Asp	Pro	Arg		Pro	Arg	Asp	Leu	
Asn	Ile	Asn	Суз		Glu	Leu	Gly	Pro		Pro	Pro	Gly	Trp	
Ile	Arg	Asn	Thr		Thr	Gly	Arg	Val		Phe	Val	Asp	His	
Asn	Arg	Thr	Thr		Phe	Thr	Asp	Pro		Leu	ser	Ala	Asn	
His	Leu	Val	Leu		Arg	Gln	Asn	Gln		Lys	Asp	Gln	Gln	
Gln	Gln	Val	Val		Leu	Cys	Pro	Asp	400 Asp	Thr	Glu	Cys	Leu	
Val	Pro	Arg	Tyr	410 Lys	Arg	Asp	Leu	Val	415 Gln	Lys	Leu	Lys	Ile	420 Leu
Arg	Gln	Glu	Leu	425 Ser	Gln	Gln	Gln	Pro	430 Gln	Ala	Gly	His	Cys	435 Arg
Ile	Glu	Val	Ser	440 Arg	Glu	Glu	Ile	Phe	445 Glu	Glu	Ser	Tyr	Arg	450 Gln
			Met	455					460			_		465
			Gly	470					475					480
_		_	Tyr	485		_		_	490	_	_			495
	_		_	500					505				_	510
			Gln	515			_		520					525
	•	-	Ser	530					535			-		540
			Arg	545		_			550					555
Ile	Asp	Gly	Gly	Phe 560	Thr	Leu	Pro	Phe	Tyr 565	Lys	Gln	Leu	Leu	Gly 570
Lys	ser	Ile	Thr		Asp	Asp	Met	Glu		Val	Asp	Pro	Asp	Leu 585
His	Asn	Ser	Leu		Trp	Ile	Leu	Glu		Asp	Ile	Thr	Gly	

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Leu Asp His Thr Phe Cys Val Glu His Asn Ala Tyr Gly Glu Ile
                605
                                     610
Ile Gln His Glu Leu Lys Pro Asn Gly Lys Ser Ile Pro Val Asn
                620
                                     625
Glu Glu Asn Lys Lys Glu Tyr Val Arg Leu Tyr Val Asn Trp Arg
                635
                                     640
                                                          645
Phe Leu Arg Gly Ile Glu Ala Gln Phe Leu Ala Leu Gln Lys Gly
                650
                                     655
                                                          660
Phe Asn Glu Val Ile Pro Gln His Leu Leu Lys Thr Phe Asp Glu
                665
                                     670
                                                          675
Lys Glu Leu Glu Leu Ile Ile Cys Gly Leu Gly Lys Ile Asp Val
                                                          690
                680
                                     685
Asn Asp Trp Lys Val Asn Thr Arg Leu Lys His Cys Thr Pro Asp
                695
                                     700
                                                          705
Ser Asn Ile Val Lys Trp Phe Trp Lys Ala Val Glu Phe Phe Asp
                710
                                     715
                                                          720
Glu Glu Arg Arg Ala Arg Leu Leu Gln Phe Val Thr Gly Ser Ser
                725
                                                          735
                                     730
Arg Val Pro Leu Gln Gly Phe Lys Ala Leu Gln Gly Ala Ala Gly
                740
                                     745
                                                          750
Pro Arg Leu Phe Thr Ile His Gln Ile Asp Ala Cys Thr Asn Asn
                755
                                     760
                                                          765
Leu Pro Lys Ala His Thr Cys Phe Asn Arg Ile Asp Ile Pro Pro
                770
                                     775
                                                          780
Tyr Glu Ser Tyr Glu Lys Leu Tyr Glu Lys Leu Leu Thr Ala Ile
                785
                                     790
Glu Glu Thr Cys Gly Phe Ala Val Glu
                800
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Thr Ile His Leu Lys Thr Leu Ile Ile Val Trp Lys Arg Tyr Ser
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                                      10
Asp Phe Lys Lys Leu His Lys Glu Leu Trp Gln Ile His Lys Asn
                 20
                                      25
Leu Phe Arg His Ser Glu Leu Phe Pro Pro Phe Ala Lys Gly Ile
Val Phe Gly Arg Phe Asp Glu Thr Val Ile Glu Glu Arg Arg Gln
                 50
                                      55
                                                           60
Tyr Ala Glu Asp Leu Leu Gln Phe Ser Ala Asn Ile Pro Ala Leu
                 65
                                      70
                                                           75
Tyr Asn Ser Lys Gln Leu Glu Asp Phe Phe Lys Gly Gly Ile Ile
                 80
                                      85
                                                           90
Asn Asp Ser Ser Glu Leu Ile Gly Pro Ala Glu Ala His Ser Asp
                 95
                                     100
Ser Leu Ile Asp Thr Phe Pro Glu Cys Ser Thr Glu Gly Phe Ser
                110
                                     115
                                                          120
Ser Asp Ser Asp Leu Val Ser Leu Thr Val Asp Val Asp Ser Leu
                125
                                     130
                                                          135
Ala Glu Leu Asp Asp Gly Met Ala Ser Asn Gln Asn Ser Pro Ile
                140
                                                          150
                                     145
Arg Thr Phe Gly Leu Asn Leu Ser Ser Asp Ser Ser Ala Leu Gly
                155
                                     160
                                                          165
Ala Val Ala Ser Asp Ser Glu Gln Ser Lys Thr Glu Glu Glu Arg
                170
                                                          180
                                     175
Glu Ser Arg Ser Leu Phe Pro Gly Ser Leu Lys Pro Lys Leu Gly
                                                          195
                185
                                     190
Lys Arg Asp Tyr Leu Glu Lys Ala Gly Glu Leu Ile Lys Leu Ala
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Leu Lys Lys Glu Glu Glu Asp Asp Tyr Glu
                                     220
<210> 409
<211> 168
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Gln Asp His Gln Phe Ala Pro Gln Ser Thr Met Ser Arg Ser Arg
                                      10
Gln Pro Pro Leu Val Thr Gly Ile Ser Pro Asn Glu Gly Ile Pro
                 20
                                      25
                                                           3.0
Trp Thr Lys Val Thr Ile Arg Gly Glu Asn Leu Gly Thr Gly Pro
                 35
                                      40
Thr Asp Leu Ile Gly Leu Thr Ile Cys Gly His Asn Cys Leu Leu
                 50
                                      55
                                                           60
Thr Ala Glu Trp Met Ser Ala Ser Lys Ile Val Cys Arg Val Gly
                                      70
                 65
Gln Ala Lys Asn Asp Lys Gly Asp Ile Ile Val Thr Thr Lys Ser
                 80
                                      85
                                                           90
Gly Gly Arg Gly Thr Ser Thr Val Ser Phe Lys Leu Leu Lys Pro
                                     100
                                                          105
                 95
Glu Lys Ile Gly Ile Leu Asp Gln Ser Ala Val Trp Val Asp Glu
                110
                                     115
                                                          120
Met Asn Tyr Tyr Asp Met Arg Thr Asp Arg Asn Lys Gly Ile Pro
                125
                                     130
                                                          135
Pro Leu Ser Leu Arg Pro Ala Asn Pro Leu Gly Met Glu Ile Glu
                                     145
                                                          150
                140
Pro Ser Thr Phe Ser Gln Lys Asp Leu Glu Met Leu Phe His Gly
                155
                                                          165
Met Ser Ala
<210> 410
<211> 108
<212> PRT
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<223> Incyte ID No: LG:407214.10.orf2:2000MAY19
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Lys Ser Arg Met Asp Leu Asp Val Val Asn Met Phe Val Ile Ala
                                      10
Gly Gly Thr Leu Ala Ile Pro Ile Leu Ala Phe Val Ala Ser Phe
                                      25
                                                           30
Leu Leu Trp Pro Ser Ala Leu Ile Arg Ile Tyr Tyr Trp Tyr Trp
                 35
                                      40
                                                           45
Arg Arg Thr Leu Gly Met Gln Val Arg Tyr Val His His Glu Asp
                 50
                                      55
                                                           60
Tyr Gln Phe Cys Tyr Ser Phe Arg Gly Arg Pro Gly His Lys Pro
                                                           75
                 65
                                      70
Ser Ile Leu Met Leu His Gly Phe Ser Gly His Lys Asp Met Trp
                 80
                                      85
Leu Ser Val Val Lys Val Pro Ser Lys Glu Pro Ala Leu Gly Leu
                 95
                                     100
Arg Gly His
<210> 411
<211> 314
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<212> PRT
<213> Homo sapiens
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<223> Incyte ID No: LG:204626.1.orf1:2000MAY19
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Gly Val Ala Arg Lys Lys Lys Ile Val Ile Lys Glu Glu Pro Lys
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Arg Arg Lys Gly Lys Met Lys Asp Arg Leu Gln Glu Leu Lys Gln
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                                                          30
Arg Thr Lys Glu Ile Glu Leu Ser Arg Asp Ser His Val Ser Thr
                                                           45
                 35
                                      40
Thr Glu Thr Glu Glu Gln Gly Val Phe Leu Gln Gln Ala Val Ile
                                                           60
                 50
                                      55
Tyr Glu Arg Glu Pro Val Ala Glu Arg His Leu His Glu Ile Gln
                                      70
                 65
Lys Leu Gln Glu Ser Ile Asn Asn Leu Ala Asp Asn Val Gln Lys
                 80
                                      85
Phe Gly Gln Gln Lys Ser Leu Val Ala Ser Met Arg Arg Phe
                                     100
                                                          105
                 95
Ser Leu Leu Lys Arg Glu Ser Thr Ile Thr Lys Glu Ile Lys Ile
                                     115
                                                          120
                110
Gln Ala Glu Tyr Ile Asn Arg Ser Leu Asn Asp Leu Val Lys Glu
                125
                                     130
                                                          135
Val Lys Lys Ser Glu Val Glu Asn Gly Pro Ser Ser Val Val Thr
                                                          150
                140
                                     145
Arg Ile Leu Lys Ser Gln His Ala Ala Met Phe Arg His Phe Gln
                155
                                     160
                                                          165
Gln Ile Met Phe Ile Tyr Asn Asp Thr Ile Ala Ala Lys Gln Glu
                                     175
                170
                                                          180
Lys Cys Lys Thr Phe Ile Leu Arg Gln Leu Glu Val Ala Gly Lys
                185
                                     190
                                                          195
Glu Met Ser Glu Glu Asp Val Asn Asp Met Leu His Gln Gly Lys
                200
                                     205
                                                          210
Trp Glu Val Phe Asn Glu Ser Leu Leu Thr Glu Ile Asn Ile Thr
                                                          225
                215
                                     220
Lys Ala Gln Leu Ser Glu Ile Glu Gln Arg His Lys Glu Leu Val
                230
                                     235
                                                          240
Asn Leu Glu Asn Gln Ile Lys Asp Leu Arg Asp Leu Phe Ile Gln
                                     250
                                                          255
                245
Ile Ser Leu Leu Val Glu Glu Gln Gly Glu Ser Ile Asn Asn Ile
                260
                                     265
                                                          270
Glu Met Thr Val Asn Ser Thr Lys Glu Tyr Val Asn Asn Thr
                                                          Lys
                275
                                     280
                                                          285
Glu Lys Phe Gly Leu Ala Val Lys Tyr Lys
                                         Lys Arg Asn Pro
                                                          Cys
                                     295
                290
                Cys Trp Cys Cys Pro Cys Cys Ser Ser Lys
Arg Val Leu Cys
                305
<210> 412
<211> 143
<212> PRT
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Met Ala Ser Glu Ser Asp Thr Glu Glu Phe Tyr Asp Ala Pro Glu
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                                                         Lys
Asp Val His Leu Gly Gly Gly Tyr Pro Val Gly Ser Pro Gly
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                 20
                                      25
Val Gly Leu Ser Thr Phe Lys Glu Thr Glu Asn Thr Ala Tyr Lys
                 35
                                      40
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Val Gly Asn Glu Ser Pro Val Gln Glu Leu Lys Gln Asp Val Ser
                 50
Asn Lys Ile Ile Glu Ser Ile Ile Glu Glu Ser Gln Lys Val Leu
                 65
Gln Leu Glu Asp Asp Ser Leu Asp Ser Thr Gly Lys Glu Leu Ser
                                                          90
                 80
                                      85
Asp Gln Ala Thr Ala Ser Pro Ile Val Ala Arg Thr Asp Leu Ser
                                                         105
                 95
                                     100
Asn Ile Pro Gly Leu Leu Ala Ile Asp Gln Val Leu Pro Glu Glu
                                                         120
                110
                                     115
Ser Gln Lys Ala Glu Ser Gln Asn Thr Phe Glu Glu Thr Glu Leu
                                                         135
                125
                                     130
Glu Phe Lys Lys Met Leu Ser Phe
                140
<210> 413
<211> 122
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
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<400> 413
Gly Arg Thr Arg Val Ser Gly Pro Val Arg Asn Asn Thr Asp Pro
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Ile His Ala Gly Arg Arg Trp Ser Ser Ser Pro Gly Arg Pro
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                                                          30
                 20
                    Trp Arg Cys Ser Cys Ser His Thr Thr
Cys Ala Arg Ser Ser
                                                          45
                 35
                                      40
Ala Gly Arg Arg Arg Trp Trp Arg Arg
                                        Pro Gly Cys Ala Trp
                 50
                                      55
                                                          60
Ala Arg Ala Ser Thr Thr Arg Ser Pro Ala Ser Pro Thr Ala Ser
                                      70
                                                           75
                 65
Ala Ala Thr Ser Ala Ser Arg Arg Thr Ala Gly Gly Pro Pro Ala
                                                          90
                 80
                                      85
Thr Ala Thr Ser Ala Thr Ala Gly Ala Arg Arg Arg Ala Lys Gln
                 95
                                     100
                                                          105
Ser Ser Ser Asn Thr Leu Gly Leu Pro Glu Leu Asn Ser Ser Ser
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                                     115
                110
Thr Lys
<210> 414
<211> 86
<212> PRT
<213> Homo sapiens
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<400> 414
Arg Pro Glu Glu Asp Met Arg Gln Leu Val His Gly Ser Gln Arg
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Gly Thr Leu Arg Lys Met Gly Leu Gln Pro Arg His Ser Ser
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                                      25
                 20
Trp Cys Gln Phe Leu Val Gly Met Val Thr Thr Phe Trp Lys Gln
                                      40
                                                           45
                 35
Gly Ala Ile Ile Ala Leu Val Ser Arg Arg Trp Lys Val Thr Arg
                                      55
                 50
                                                           60
Lys Gly Trp Gln Cys Gln Val Arg Thr Thr Leu Ala Cys Arg Leu
                 65
                                      70
Leu Asp Cys Ile Leu Pro Pro Asn Ser Tyr Asn
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85

80

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<211> 213
<212> PRT
<213> Homo sapiens
<220>
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Arg Gln Trp Gln Leu Leu Cys Cys Pro Cys Cys Cys Gly Gly Cys
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Trp Ala Arg Val Glu Val Gln Gly Trp Gly His Gly Leu Ala Pro
                 20
Gly Leu Gly Trp Ser Cys Pro Ala Gly Pro Gly Gly His Arg Gly
                                                           45
                 35
                                      40
Cys Gly Cys Pro Gly Val Gly Arg Thr Trp Lys Ala Ala Arg Ser
                                      55
                                                           60
Arg Gly Ser Trp Val Ser Trp Val Ser Trp Gly Thr Trp Arg Ser
                                                          75
                 65
                                      70
Trp Arg Ser Trp Gly Ser Trp Gly Ser Trp Asp Ser Arg Ala Pro
                                      85
                                                          90
                 80
Pro Pro Glu Pro Ser Arg Thr Pro Arg Gly Thr Asn Arg Ser Arg
                 95
                                                          105
                                     100
Thr Ser Pro Pro Ala Cys Arg Arg Ala Ser Arg Ser Gly Pro Arg
                                     115
                                                          120
                110
Ala Pro Thr Pro Pro Pro Arg Pro Gln Arg Arg Ser Ala Ala Ala
                125
                                     130
                                                          135
Cys Gly Trp Arg Pro Arg Ser Gly Arg Pro Gly Arg Gly Leu Ser
                140
                                     145
                                                          150
Trp Arg Pro Arg Arg Pro Gly Ala Arg Gln Arg Ser Trp Ser
                                                          165
                155
                                     160
                    Cys Arg Cys Arg Ala Arg Arg Pro Thr Ala
Ala Ser Phe Gly Arg
                                     175
                                                          180
                170
Cys Gly Ala Gly Ser Asp Gly Leu Ala Leu Ala Gly Thr Arg Ala
                                     190
                                                          195
                185
Ser Arg Arg Ala Arg Cys Arg Arg Ser Arg Ser Arg Ala Gly Cys
                200
                                     205
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Gly Gly Ser
<210> 416
<211> 263
<212>, PRT
<213> Homo sapiens
<220>
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<223> Incyte ID No: LG:482261.1.orf1:2000FEB18
<400> 416
Leu Leu Asn Phe Val Ser Leu Ser Leu Phe Leu Phe Ser Phe Pro
 1
                                      10
                                                           15
Arg Leu His Arg Glu Gly Glu Ser Leu Arg Val Val Tyr Gln Ala
                 20
                                      25
                                                           3.0
Gly Ser Pro Thr Ser Leu Ala Pro Asn Thr Val Ser Ser Asn Pro
                 35
                                      40
                                                           45
                                                          Va1
Gly Glu Val Thr Pro Glu Arg Gly Arg Cys Glu Glu Arg Ser
                                                           60
                 50
                                      55
Gln Glu Leu Pro Arg Thr Cys Gly Arg Pro Cys Gly Lys Leu Val
                                                           75
                 65
                                      70
His Ser Glu His Ser Arg Asp Thr Met Gly Gln Ser Lys
                                                     Ser Lys
                 80
                                      85
His Ser Ala Tyr Leu His Phe Ile Lys Leu Leu Lys Arg Ala
                 95
                                     100
Gly Ile Lys Ala Ser Thr Glu Asn Leu Ile Thr Leu Phe Pro Thr
                110
                                     115
```

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Val Glu Gln Tyr Cys Pro Trp Phe Pro Glu His Gly Thr Met Asp
                125
                                     130
                                                          135
Phe Lys Asp Trp Glu Gln Val Gly Ile Ala Leu Lys Gln Val Cys
                140
                                     145
                                                          150
Lys Glu Gly Lys Phe Ile Pro Leu Thr Ala Trp Ser Asn Trp Ala
                155
                                     160
                                                          165
Ile Val Lys Ala Ala Ser Glu Pro Phe Gln Ser Glu Asn Glu Ala
                170
                                     175
                                                          180
Tyr Pro Pro Ala Glu Arg Ile Ser Ala Glu Glu Gly Gly Asp Ala
                185
                                     190
                                                          195
Ala Glu Gly Gly Glu Asp Ser Glu Glu Asp Phe Glu Glu Asn Thr
                200
                                     205
                                                          210
Asp Lys Pro Gly Asp Glu Leu Ile Ser Phe Glu Glu His Val Gly
                215
                                     220
                                                          225
Pro Ser Ala Ala Pro Lys Ile Glu Lys Pro Tyr Met Pro Arg
                                                          Cys
                                     235
                                                          240
                230
Leu Lys Gln Arg Arg Ala Leu Arg Ser Ser Arg Leu Leu Ile
                                                          Gly
                                                          255
                245
                                     250
Ile Ile Arg Ser Gly Arg Leu Gln
                260
<210> 417
<211> 175
<212> PRT
<213> Homo sapiens
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<223> Incyte ID No: LG:480328.1.orf1:2000FEB18
<400> 417
Val Arg Cys Gly Gly Gln Ser Gly Ser Arg Cys Glu Arg Val Pro
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Gln Lys Lys Thr Trp Leu Pro Lys Cys Leu Ser Pro Ser Ala Ser
                                      25
                                                           30
                 20
Leu Gly Leu Ala Leu Ala Val Ala Gly Gly Met Val Asn Ser Ala
                 35
                                      40
Leu Cys Asn Val Asp Ala Gly His Arg Ala Ala Ile Phe Asp Gln
                                      55
Phe Arg Gly Val Gln Asn Ile Val Val Gly Glu Gly Thr His Phe
                 65
                                      70
                                                           75
Leu Ile Pro Cys Val Gln Lys Pro Ile Ile Phe Asp Cys Cys Ser
                                                           90
                 80
                                      85
Gln Pro Arg Ser Ala Pro Val Ile Thr Gly Ser Lys Asp Leu Gln
                                     100
                                                          105
                 95
Asn Val Asn Ile Thr Leu Cys Ile Leu Phe Arg Pro Ile Thr Ser
                110
                                     115
                                                          120
Gln Leu Pro Arg Ile Phe Thr Ser Ile Gly Glu Asp Tyr Asp Glu
                125
                                     130
                                                          135
Cys Val Leu Pro Phe Ile Thr Thr Glu Ile Leu Lys Ser Leu Val
                140
                                     145
                                                          150
Ala Arg Phe Asp Ala Gly Glu Leu Ile Thr Gln Arg Glu Leu Val
                155
                                     160
                                                          165
Ser Ser Gln Val Ser Asn Asn Leu Met Glu
                170
<210> 418
<211> 272
<212> PRT
<213> Homo sapiens
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<400> 418
Gln Val Pro Phe Pro Ser His Ile Pro Ile Tyr Val Cys Leu Cys
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Val Gln Gln Thr Pro Ala Phe Ala Thr Met Leu Ser Ser Thr Asp
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Phe Thr Phe Ala Ser Trp Glu Leu Val Val Arg Val Asp His Pro
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Asn Glu Glu Gln Gln Lys Asp Val Thr Leu Arg Val Ser Gly Asp
                 50
                                      55
                                                           60
Leu His Val Gly Gly Val Met Leu Lys Leu Val Glu Gln Ile Asn
                                                          75
                 65
                                      70
Ile Ser Gln Asp Trp Ser Asp Phe Ala Leu Trp Trp Glu Gln Lys
                                      85
                 80
His Cys Trp Leu Leu Lys Thr His Trp Thr Leu Asp Lys Tyr Gly
                                     100
                 95
Val Gln Ala Asp Ala Lys Leu Leu Phe Thr Pro Gln His Lys Met
                110
                                     115
Leu Arg Leu Arg Leu Pro Asn Leu Lys Met Val Arg Leu Arg Val
                                     130
                                                          135
                125
Ser Phe Ser Ala Val Val Phe Lys Ala Val Ser Asp Ile Cys Lys
                                                          150
                140
Ile Leu Asn Ile Arg Arg Ser Glu Glu Leu Ser Leu Leu Lys Pro
                155
                                     160
                                                          165
Ser Gly Asp Tyr Phe Lys Lys Lys Lys Lys Asp Lys Asn Asn
                170
                                     175
                                                         180
Lys Glu Pro Ile Ile Glu Asp Ile Leu Asn Leu Glu Ser Ser Pro
                                                          195
                                     190
                185
Thr Ala Ser Gly Ser Ser Val Ser Pro Gly Leu Tyr Ser Lys Thr
                                     205
                200
Met Thr Pro Ile Tyr Asp Pro Ile Asn Gly Thr Pro Ala Ser Ser
                215
                                     220
                                                          225
Thr Met Thr Trp Phe Ser Asp Ser Pro Leu Thr Glu Gln Asn Cys
                230
                                     235
                                                          240
Ser Ile Leu Ala Phe Ser Gln Pro Pro Gln Ser Pro Glu Ala Leu
                                     250
                245
                                                          255
Ala Asp Met Tyr Gln Pro Arg Ser Leu Val Asp Thr Ala Lys Leu
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Asn Ala
<210> 419
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<223> unknown or other
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Arg Xaa Arg Gln Pro Leu Leu Gln Ser His Pro Glu Ala Asp Trp
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Ser Thr His Gly Arg Ser Met Arg Lys Leu Ile Val Arg Phe Ile
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Phe Leu Lys Phe Trp Thr Tyr Thr Val Arg Ala Ser Thr Asn Leu
Thr Gln Asn Gly Asp Cys Ser Gln Cys Ile Tyr Gln Val Thr Glu
                 50
                                      55
                                                           60
Val Gly Gln Gln Ile Lys Thr Ile Phe Leu Phe Tyr Ser Tyr Tyr
                  65
                                      70
Glu Cys Met Glu Thr Leu Lys Glu Thr Cys Leu Tyr Asn Ala Thr
                 80
                                      85
Gln Tyr Lys Val Cys Ser Pro Arg Asn Asp Arg Pro Asp Ala Cys
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                                     100
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Tyr Asn Pro Ser Glu Pro Ala Ala Thr Thr Val Phe Glu Ile Arg
                110
                                     115
                                                         120
Thr Gly Leu Leu Gly Asp Thr Ser Lys Ile Ile Thr Arg Thr
                125
                                     130
                                                         135
Glu Glu Lys Glu Ile Pro Lys Gln Ile Thr Leu Arg Phe Asp Ala
                                                         150
                140
                                     145
Cys Ala Ala Ile Asn Ser Lys Lys Leu Glu Ile Gly Cys Gly
                                     160
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                155
Leu Asn
<210> 420
<211> 59
<212> PRT
<213> Homo sapiens
<220>
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<223> Incyte ID No: LG:399395.1.orf2:2000MAY19
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Ser Gln His Phe Gly Arg Pro Arg Gln Glu Asp His Leu Ser Pro
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Gly Val Gln Asp Gln Pro Gly Gln His Ser Glu Thr Leu Thr Gln
                                      25
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Lys Ile Lys Arg Lys Asp Lys Asn Thr Arg Met Ala Lys Gln Thr
                 35
                                      40
Ser Val His Gln Pro Gly Gly Ile Leu Tyr Ser Leu Leu Lys
<210> 421
<211> 216
<212> PRT
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<220>
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<400> 421
Ser Pro Arg Pro Leu Gln Ser Ala Gly Glu Gly Val Thr His Val
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Leu Ile Leu Leu Glu Ser Pro Ala Arg Pro Val Ala Ala Val Thr
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Gln Val Gln Arg Arg Arg Tyr His Arg Leu Ser Asp Met Ser Met
                                                          45
                                      40
Leu Ala Glu Arg Arg Lys Gln Lys Trp Ala Val Asp Pro Gln
                                      55
                                                          60
                 50
Asn Thr Ala Trp Ser Asn Asp Asp Ser Lys Phe Gly Gln Arg Met
                 65
                                      70
                                                          75
Leu Glu Lys Met Gly Trp Ser Lys Gly Lys Gly Leu Gly Ala Gln
                 80
                                      85
                                                          90
Glu Gln Gly Ala Thr Asp His Ile Lys Val Gln Val Lys Asn Asn
                                                          105
                                     100
                 95
His Leu Gly Leu Gly Ala Thr Ile Asn Asn Glu Asp Asn Trp Ile
                                                          120
                110
                                     115
Ala His Gln Asp Asp Phe Asn Gln Leu Leu Ala Glu Leu Asn Thr
                                     130
                125
Cys His Gly Gln Glu Thr Thr Asp Ser Ser Asp Lys Lys Glu Lys
                                     145
                                                          150
                140
Lys Ser Phe Ser Leu Glu Glu Lys Ser Lys Ile Ser Lys Asn Arg
                155
                                     160
                                                          165
Val His Tyr Met Lys Phe Thr Lys Gly Arg Cys Gln Ser Leu His
                170
                                     175
                                                         180
Ser Arg Gly Glu Arg Asn His Asp Asn Gln Arg Leu His His Pro
                                     190
                185
Gly Val Leu Cys Gln Ala Asp Gly Ser Thr Glu Glu Gln Ala Pro
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210
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Gly Ser Ser Ser Arg Val
<210> 422
<211> 162
<212> PRT
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Ser Ala Thr Ala Ala Ala Pro Arg Glu Arg Lys Met Ala Pro His
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Gly Pro Gly Ser Leu Thr Thr Leu Val Pro Trp Ala Ala Ala Leu
                                      25
                - 20
Leu Leu Ala Leu Gly Val Glu Arg Ala Leu Ala Leu Pro Glu Ile
                 35
                                      40
Cys Thr Gln Cys Pro Gly Ser Val Gln Asn Leu Ser Lys Val Ala
                 50
                                      55
                                                           60
Phe Tyr Cys Lys Thr Thr Arg Glu Leu Met Leu His Ala Arg Cys
                                                           75
                 65
                                      70
Cys Leu Asn Gln Lys Asp Pro Gly Pro Asn Phe His Gln Ala His
                                                           90
                 80
                                      85
Thr Thr Val Ile Ile Asp Leu Gln Ala Asn Pro Leu Lys Gly Asp
                                     100
                                                          105
                 95
Leu Ala Asn Thr Phe Arg Gly Phe Thr Gln Leu Gln Thr Leu Ile
                                                          120
                                     115
                110
Leu Pro Gln His Val Asn Cys Ser Gly Gly Ile Asn Ala Trp Asn
                125
                                     130
                                                          135
Thr Ile Thr Ser Tyr Ile Asp Asn Gln Ile Cys Gln Gly Gln Lys
                140
                                     145
Asn Leu Cys Asn Asn Thr Gly Asp Pro Glu Met Cys
                155
                                     160
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